# Production and Isolation of Two Novel Esperamicins in a Chemically Defined Medium

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Esperamicins, a family of extremely potent compounds, showing a broad spectrum of antimicrobial and antitumor activities in murine models has been identified in cultures of *Actinomadura verrucosospora* ATCC 39334<sup>1,2)</sup>. The isolation and structure elucidation of esperamicins  $A_1$  (esp  $A_1$ ),  $A_{1b}$  (esp  $A_{1b}$ ) and  $A_2$  (esp  $A_2$ ) have been reported<sup>1,3,4)</sup>. The esperamicins consist of a bicyclic core to which attached a trisaccharide and a substituted 2-deoxy-L-fucose. The bicyclic core contains an enediyne, an allylic trisulfide and a bridgehead eneone.

Esp  $A_1$  (Fig. 1), the major component of the esperamicin family isolated from the fermentation of *A. verrucosospora* ATCC 39334 grown in the complex medium H946<sup>5)</sup>, is one of the most potent antitumor agents yet discovered<sup>2)</sup>. Three minor esperamicins, esp  $A_2$ , esp  $A_{1b}$  and esp P (Fig. 1) were also isolated from the large scale fermentation of *A. verrucosospora* ATCC 39334 grown in the complex medium<sup>5,6)</sup>. While developing a defined medium for more efficient incorporation of labelled precursors into esp  $A_1$  during the biosynthetic studies<sup>7)</sup>, we detected the formation of two new esperamicin analogs in a defined medium. We report here the production, isolation, structure determination and antitumor activity of these analogs.

A vegetative culture of A. verrucosospora ATCC 39334 was prepared by transferring 4 ml of the frozen stock culture to a 500-ml Erlenmeyer flask containing 100 ml of a vegetative medium consisting of starch 2%, glucose 0.5%, Pharmamedia 1%, yeast extract 1% and CaCO<sub>3</sub> 0.2%. The vegetative culture was incubated at 28°C for 96 hours on a rotary shaker operating at 250 rpm. An 8-ml aliquot of incubated seed culture was transferred to a 500-ml Erlenmeyer flask containing 100 ml of complex medium H946. Medium H946 was prepared using cane molasses 6%, starch 2%, fish meal 2%, CuSO<sub>4</sub> · 5H<sub>2</sub>O 0.01%, CaCO<sub>3</sub> 0.2% and NaI 0.00005%. The metabolite production profile of an extract from a 10-day old culture of A. verrucosospora ATCC 39334 grown in the complex medium H946 at 28°C on a rotary shaker operating at 250 rpm analyzed by HPLC is shown in Fig. 2. HPLC analysis was carried out by using a C-18 reverse-phase column (Novapak, 3.9 × 150 mm, Waters Associates), the solvent system was 0.05 M ammonium acetate (pH 4.5)-CH<sub>3</sub>OH-CH<sub>3</sub>CN (1:1:1) at a flow rate of 1 ml/minute and detected at 254 nm. The major product of fermentation was esp A<sub>1</sub> (Rt. 8.6 minutes) which comprised 75% of the esperamicin complex. The other major component of the fermentation,  $esp A_2$  (Rt. 16.3 minutes) constituted approximately 15% of the complex.

Using the same medium and conditions described above for the preparation of the seed culture, an 8-ml aliquot of incubated seed culture was transferred to a

#### Fig. 1. Naturally occuring esperamicins.

	Сн₃			CH <sub>3</sub> 0 0 CH <sub>3</sub> O <sub>2</sub> (	
					$OR_2$
Esperamicin	n	R	R <sub>1</sub>	R <sub>2</sub>	
Esperamicin A <sub>1</sub>	n 3	R CH(CH <sub>3</sub> ) <sub>2</sub>	R <sub>1</sub> AC	R <sub>2</sub> H	$AC: \qquad O \qquad H \qquad CH_2$
$\frac{\text{Esperamicin}}{A_1}$	n 3 3	R CH(CH <sub>3</sub> ) <sub>2</sub> CH <sub>2</sub> CH <sub>3</sub>	R <sub>1</sub> AC AC	R <sub>2</sub> H H	$AC: \qquad H \qquad CH_2 \\ H \qquad H \qquad H \qquad OCI$
$\frac{A_1}{A_{1b}}$	n 3 3 3	R CH(CH <sub>3</sub> ) <sub>2</sub> CH <sub>2</sub> CH <sub>3</sub> CH <sub>3</sub>	R <sub>1</sub> AC AC AC	R <sub>2</sub> H H H	$AC: \qquad H \qquad CH_2 \\ H \qquad H \qquad OCI$
Esperamicin $A_1$ $A_{1b}$ $A_{1c}$ P	n 3 3 3 4	R CH(CH <sub>3</sub> ) <sub>2</sub> CH <sub>2</sub> CH <sub>3</sub> CH <sub>3</sub> CH(CH <sub>3</sub> ) <sub>2</sub>	R <sub>1</sub> AC AC AC AC	R <sub>2</sub> H H H H	$AC: \qquad 0 \qquad H \qquad CH_2$
Esperamicin $A_1$ $A_{1b}$ $A_{1c}$ P $A_2$	n 3 3 3 4 3	$\begin{array}{c} R\\ \hline CH(CH_3)_2\\ CH_2CH_3\\ CH_3\\ CH_3\\ CH(CH_3)_2\\ CH(CH_3)_2 \end{array}$	R <sub>1</sub> AC AC AC AC H	R <sub>2</sub> H H H H AC	$AC: \qquad OH \qquad CH_2$ $H \qquad H \qquad OH \qquad OH$ $CH_3O \qquad H \qquad O$

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500-ml Erlenmeyer flask containing 100 ml of a chemically defined medium DF-15. Shifting the fermentation of A. verrucosospora ATCC 39334 from the complex medium to the defined medium DF-15 using sucrose as the sole carbon source and ammonium sulfate as the sole nitrogen source yielded a significantly different production profile for the esperamicin complex. Defined medium DF-15 consisted of sucrose 4%, NH<sub>4</sub>Cl 0.2%, Na<sub>2</sub>SO<sub>4</sub> 0.2%, K<sub>2</sub>HPO<sub>4</sub> 0.1%, MgCl<sub>2</sub>·6H<sub>2</sub>O 0.1%, NaCl 0.1%, CaCO<sub>3</sub> 0.2%, MnCl<sub>2</sub>·4H<sub>2</sub>O 0.0001%, ZnCl<sub>2</sub> 0.0001%, FeCl<sub>2</sub>·4H<sub>2</sub>O 0.0001% and NaI 0.00005%. The metabolite production profile of an extract from a 6-day old culture of A. verrucosospora ATCC 39334 grown in medium DF-15 analyzed by HPLC is shown in Fig. 3. Esp  $A_1$  and esp  $A_2$  were not detected in the extract of A. verrucosospora ATCC 39334 grown in medium DF-15, however, two new esperamicin metabolites (Rts. 6.5





Fig. 3. Chromatogram of HPLC analysis of an extract from a 6-day old culture of *Actinomadura verrucosospora* ATCC 39334 grown in defined medium DF-15.



minutes and 10.3 minutes) that had not been previously isolated in our laboratory were identified by HPLC analysis.

The purification of the two new analogs from a 10-liter fermentation broth is summarized in Fig. 4. The structure determination of these two new analogs is summarized in Fig. 5. Mass spectrometry analysis shows that both analogs have identical molecular ions 28 mass unit less than that of esp  $A_1$  indicating a difference of two methylene groups. In addition, fragmentation of the glycosidic bond of the aminosugar residue results in formation of base peaks at m/z 144 that clearly identifies the alkylation pattern of the analyzed congeners (Fig. 5). Lack of availability in sufficient quantities, their instability, and the complexity of their NMR spectra precluded a full assignment of the chemical shifts for the minor components, yet the close structural relations among them allowed us to classify them based on a few readily identifiable diagnostic peaks. In the <sup>1</sup>H spectra the lowest three singlets at 11.77 ppm (br, NH), 8.58 ppm (H-12<sup>iv</sup>), and 7.47 ppm (H-9<sup>iv</sup>) are indicative of the  $A_1$ series, while the equivalent peaks for the A<sub>2</sub> series are found at 11.90, 8.61, and 7.57 ppm, respectively (Fig. 5). In the high field region of the esp  $A_{1c}$  spectrum a doublet at 1.30 ppm indicates the presence of  $6^{iv}$  methyl group. The equivalent doublet for the A2c congener was shifted to 1.24 ppm (Fig. 5). Additional diagnostic information can be obtained from <sup>13</sup>C NMR data of the methyl region of the spectra. The signals for the N-isopropyl groups in esp  $A_1$  and esp  $A_2$  found at 22.2, 22.3 and 47.2 ppm were replaced by the N-methyl signals at 34.1 and 33.6 ppm for esp  $A_{1c}$  and esp  $A_{2c}$ , respectively (Fig. 5). The above data clearly indicate that the two new esperamicins were esp  $A_{1c}$  and esp  $A_{2c}$ . Esp  $A_{1c}$  was later determined to be the same as FR-900406 reported by Fujisawa Pharm. Co. Ltd.<sup>8,9)</sup>. Esp A<sub>2c</sub> is a novel esperamicin analog that has not been previously reported.



Fig. 4. Scheme for isolation of esperamicins  $A_{1c}$  and  $A_{2c}$  from a 10 liter shake flask batch.

The extracts and fractions from purification were evaporated to dryness *in vacuo* using a rotary evaporator at 25°C. Fig. 5. Structure determination of esperamicins A<sub>1e</sub> and A<sub>2e</sub>. HPLC separation conditions used in LC-MS study: C-18 Novapak column (3.9×150 mm, Waters Associates); CH<sub>3</sub>CN-CH<sub>3</sub>OH-H<sub>2</sub>O (33:33:34) solvent system; 1 ml/minute flowrate; temperature; 25°C.



Esperamicin		$A_{1c} (R_1 = R, R_2 = H)$	$A_{2c} (R_1 = H, R_2 = R)$
HPLC			
Rt, (min):		6.50	16.10
MS			
LC-TSP $(m/z)$ :	$(M + H)^{+}$	1297	1297
	$(M + Na)^+$	1319	1319
	Base peak	144	144
NMR		4	
$^{1}$ H, CDCl <sub>3</sub> (ppm):	13 <sup>1V</sup> -NH	11.77	11.90
, , ,	12 <sup>IV</sup> -H	8.58	8.61
	9 <sup>rv</sup> -H	7.47	7.56
	6 <sup>IV</sup> -H	1.30	1.24
$^{13}C$ , CDCl <sub>3</sub> (ppm):	C-4'''-SMe	13.7	13.7
, j.u.,	C-6 <sup>IV</sup>	16.7	16.6
	C-6′	17.6	17.5
	C-15-S <sub>3</sub> Me	22.7	22.7
	C-4″-NH <i>Me</i>	34.1	33.6

Table 1. Effect of esperamicins  $A_1$ ,  $A_{1e}$  and  $A_{2e}$  on P388 leukemia.

Compound	Dose (mg/kg/inj, ip)	Effect (% T/C)	Compound	Dose (mg/kg/inj, ip)	Effect (% T/C)
Esp A <sub>1</sub>	0.016	180		0.001	135
	0.008	155		0.0005	130
	0.004	160	Esp A <sub>2s</sub>	0.016	145
	0.002	135		0.008	135
	0.001	135		0.004	125
	0.0005	130		0.002	110
Esp A <sub>10</sub>	0.016	175		0.001	110
1 10	0.008	160		0.0005	105
	0.004	150	Control	0	100
	0.002	155			

Tumor inoculum: 10<sup>6</sup> ascites cells, ip.

Host: CDF<sub>1</sub> mice.

Schedule:  $Q1D \times 1$ ; 2.

Evaluation: Medium survival time (MST).

Effect: % T/C = (MST treated/MST control)  $\times$  100.

Criteria: % T/C > 125 considered significant antitumor activity.

Control mice were given saline injection.

Compound	Dose (mg/kg/inj, ip)	Effect (% T/C)	Compound	Dose (mg/kg/inj, ip)	Effect (% T/C)
Esp A <sub>1</sub>	0.01	56		0.01	144
	0.003	104		0.003	144
	0.001	152		0.001	144
	0.0003	152		0.0003	136
	0.0001	144		0.0001	128
	0.00003	128		0.00003	120
	0.00001	128		0.00001	120
Esp $A_2$	0.1	48	Control	0	100
	0.03	104			

Table 2. Effect of esperamicins  $A_1$  and  $A_2$  on P388 leukemia.

Tumor inoculum: 10<sup>6</sup> ascites cells, ip.

Host: CDF<sub>1</sub> mice.

Schedule:  $Q1D \times 9$ ; 1.

Evaluation: Medium survival time (MST).

Effect: % T/C = (MST treated/MST control)  $\times$  100.

Criteria: % T/C > 125 considered significant antitumor activity.

Control mice were given saline injection.

The *in vivo* antitumor activity of esp  $A_1$ , esp  $A_{1c}$  and esp  $A_{2c}$  against P388 leukemia implanted ip in mice is shown in Table 1. The activity and potency of esp  $A_{1e}$ against P388 leukemia are similar to those of esp  $A_1$  and both of the compounds were active at the lowest dosage tested, 0.0005 mg/kg/inj. Esp A<sub>2c</sub> was active against P388 leukemia at 0.004 mg/kg/inj but was inactive at 0.002 mg/kg/inj, indicating that it is at least 8 fold less potent than  $\exp A_1$  and  $\exp A_{1c}$  based on minimal effective dose comparisons. Table 2 compares the effect of esp  $A_1$  and esp  $A_2$  against P388 leukemia implanted ip in mice. Esp  $A_1$  was again active at the lowest dosage tested, 0.00001 mg/kg/inj. Esp A2 was active against P388 leukemia at 0.0001 mg/kg/inj but was inactive at 0.00003 mg/kg/inj, indicating that it is at least 10 fold less potent than esp A<sub>1</sub> based on minimium effective dose comparisons. The above finding indicates that the position of attachment of the anthranilate chromophore to the 2-deoxy-L-fucose moiety is an important factor in determining the antitumor potency of esperamicins. Mechanism of action studies<sup>10)</sup> demonstrated that the anthranilate-deoxyfucose moiety affects the uptake of the drug into the cells and after activation may also sterically hinder the resulting phenylene diradical so as to lead to inefficient double stranded breakage of DNA. Furthermore, results from molecular dynamics simulations and spectroscopic studies<sup>11~13</sup> showed that the anthranilatedeoxyfucose moiety contributes significantly to the biochemistry of esp A<sub>1</sub>-induced DNA damage.

In this paper, we have demonstrated that new esperamicin analogs can be generated by changing the composition of the fermentation medium. The comparison of antitumor activity between esp  $A_{1c}$  and esp  $A_{2c}$  also confirms that the  $A_1$  series are more potent than the  $A_2$  series of esperamicin in the *in vivo* antitumor tests.

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