

NEW ANTHRACYCLINE  
ANTIBIOTICS, RHODIRUBINS

Sir:

We have previously reported on aclacinomycins A and B<sup>1)</sup> and will report on their nineteen analogues<sup>2)</sup> produced by a strain No. MA144-M1, which was classified as *Streptomyces galilaeus*. In a continuing search for other new anthracycline antibiotics, we have found rhodirubins in the culture broths of a strain designated as ME505-HE1 belonging to *Streptomyces* sp. In this communication, the isolation of rhodirubins and their properties are reported.

The strain ME505-HE1 was cultivated at 28°C for 5 days on a rotary shaker in a medium consisting of 2% potato starch, 2% glucose, 2% soybean meal, 0.5% yeast extract, 0.25% NaCl, 0.3% CaCO<sub>3</sub>, 0.0003% CuSO<sub>4</sub>·5H<sub>2</sub>O, 0.0003% MnCl<sub>2</sub>·4H<sub>2</sub>O and 0.0003% ZnSO<sub>4</sub>·7H<sub>2</sub>O, pH 7.4. Red pigments in the mycelial cake were extracted with acetone, evaporated under reduced pressure and re-extracted with chloroform. After concentration, the dried residue was dissolved in a small amount of ethyl acetate and crude rhodirubin was precipitated by addition of 10 times volume of *n*-hexane. The crude dark red powder thus obtained was dissolved in a small amount of toluene and subjected to a silicic acid column chromatography. After eluting the red non-carbohydrate containing pigments with toluene-methanol in 100:2 volume ratio, rhodirubins C, G, B and A were successively eluted with a 100:3 mixture of the same solvents. Rhodirubins D and E were eluted with a 10:1 mixture. Each rhodirubin fraction described above was subjected to preparative silicic acid thin-layer chromatography using chloroform-methanol (20:1 or 10:1). Thus, 33 mg of rhodirubin A, 87.6 mg of rhodirubin B and 105 mg of rhodirubin C were obtained in pure red amorphous states from 2.4 g of the crude powder described above. The minor components D, E and G were obtained in quantities sufficient only for determination of TLC Rf values and for analysis of sugar moieties by acid hydrolysis. Their Rf values on a silicic acid thin-layer chromatography (Kieselgel 60F<sub>254</sub>, Merck Co.) with several solvent systems are shown in Table 1.

On acid hydrolysis in 0.1 N HCl for 30 minutes at 85°C, all rhodirubins gave  $\epsilon$ -pyrromycinone<sup>3,4)</sup>. Thin-layer chromatography of acid hydrolysates

with *n*-butanol-acetic acid-water (4:1:1) indicated that rhodirubin A contained three hexoses corresponding to rhodosamine, 2-deoxyfucose and rhodinos, B contained rhodosamine and two moles of rhodinos, C contained rhodosamine, 2-deoxyfucose and cinerulose, E contained rhodosamine and 2 moles of 2-deoxyfucose, and G contained rhodosamine, 2-deoxyfucose and D-cinerulose, while D had two hexoses corresponding to rhodosamine and 2-deoxyfucose, as shown in Table 2. For identification of the hexoses, authentic samples of aclacinomycin A<sup>1)</sup>, MA144 G1<sup>2)</sup> and streptolydigin<sup>5)</sup> were hydrolyzed and the hexoses thus obtained were compared with those obtained from hydrolysis of rhodirubins. Rhodirubin C was found to be identical with cinerubin A<sup>6)</sup> by the mixed melting point determination, IR, PMR and <sup>13</sup>C-NMR spectra, and the physicochemical properties of the methyl disaccharide obtained by their methanolysis in methanol containing 0.01 N HCl for 30 minutes.

Physicochemical properties of rhodirubins A and B are as follows: Rhodirubin A: m.p. 141~143°C;  $[\alpha]_D^{20} + 120^\circ$  (*c* 1.0, CHCl<sub>3</sub>);

Anal. calcd. for C<sub>42</sub>H<sub>55</sub>NO<sub>16</sub>: C 60.77, H 6.67, O 30.81, N 1.69.

Found: C 60.39, H 6.63, O 30.72, N 1.71.

The ultraviolet and visible absorption spectra in methanol show the following maxima ( $E_{1\%}^{1\text{cm}}$ ): 234 (527), 258 (280), 292 (120), 412sh (53), 465sh (133), 480 (153), 490 (160), 510sh (127), 524sh (107), 570 nm (20). The IR spectrum in KBr shows peaks at the following wavelengths in cm<sup>-1</sup>: 3430, 2950, 2930, 2810, 2750, 1735, 1640, 1600, 1450, 1320, 1300, 1220, 1160, 1120, 1040, 1000, 970, 960, 920, 800 and 760. The PMR spectrum of rhodirubin A in CDCl<sub>3</sub> (100 MHz)

Table 1. Rf values of rhodirubins

Components	Solvent systems		
	(1)	(2)	(3)
Rhodirubin A	0.52	0.48	0.26
Rhodirubin B	0.52	0.56	0.29
Rhodirubin C	0.88	0.70	0.53
Rhodirubin D	0.28	0.26	0.13
Rhodirubin E	0.28	0.17	0.11
Rhodirubin G	0.85	0.66	0.49

Solvent systems:

(1) chloroform-methanol (10:1)

(2) chloroform-methanol-formic acid (9:1:0.5)

(3) benzene-ethyl acetate-methanol (5:5:1)

Table 2. Rf values of hexoses obtained by hydrolysis of rhodirubins

Rhodirubins	Rhodosamine Rf=0.14	2-Deoxy-fucose 0.55	Rhodi-nose 0.70	Cineru-lose 0.81
A	+	+	+	-
B	+	-	++	-
C	+	+	-	+
D	+	+	-	-
E	+	++	-	-
G	+	+	-	+

Rf: Silica gel thin-layer chromatography with *n*-butanol - acetic acid - water (4: 1: 1)

Table 3. Antimicrobial spectrum of rhodirubins A and B

Organisms	MIC, $\mu\text{g/ml}$	
	Rhodiru-bin A	Rhodiru-bin B
<i>Bacillus subtilis</i> ATCC6633	0.78	1.56
<i>Bacillus cereus</i> ATCC9634	0.2	0.4
<i>Bacillus megaterium</i> NRRL B-938	0.78	1.56
<i>Staphylococcus aureus</i> FDA209P	1.56	1.56
<i>Staphylococcus aureus</i> Smith	0.4	0.78
<i>Sarcina lutea</i> ATCC9341	0.4	0.78
<i>Micrococcus flavus</i>	<0.2	<0.2
<i>Corynebacterium bovis</i> 1810	0.2	0.4
<i>Pseudomonas fluorescens</i> NIHJ B254	100	100
<i>Proteus morganii</i>	>100	>100
<i>Mycobacterium smegmatis</i> ATCC607	6.25	3.1
<i>Candida albicans</i> IAM4905	>100	>100
<i>Candida tropicalis</i> IAM4942	>100	>100

shows the following chemical shifts (ppm): 7.6, s; 7.24, s; 5.50, m; 5.62, m; 5.02, m; 4.84, m; 4.52, q; 4.7 ~ 3.90, overlapping m; 3.72, s; 3.60 ~ 0.09, overlapping m and 2.18 s. Rhodirubin B: m.p. 135 ~ 137°C;  $[\alpha]_D^{20} + 190^\circ$  (c 0.1,  $\text{CHCl}_3$ );

Anal. calcd. for  $\text{C}_{42}\text{H}_{55}\text{NO}_{15}$ : C 61.99, H 6.77, O 29.52, N 1.72.

Found: C 61.23, H 6.80, O 28.77, N 1.94.

The ultraviolet and visible absorption spectra in methanol show the following maxima ( $E_{1\%}^{1\text{cm}}$ ): 235 (600), 257 (310), 290 (120), 410sh (53), 464sh (153), 476sh (173), 491 (187), 510sh (147), 522sh (113) and 570 nm (19). The infrared spectrum in KBr shows peaks at the following wavelengths in  $\text{cm}^{-1}$ : 3470, 2960, 2940, 2820, 2790, 1740, 1640, 1600, 1450, 1300, 1220, 1160, 1120, 1040,

Fig. 1. Structure of rhodirubins A and B

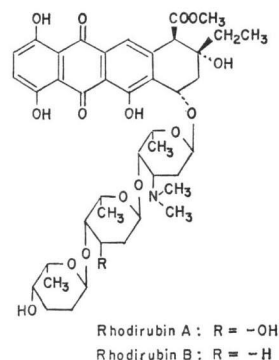
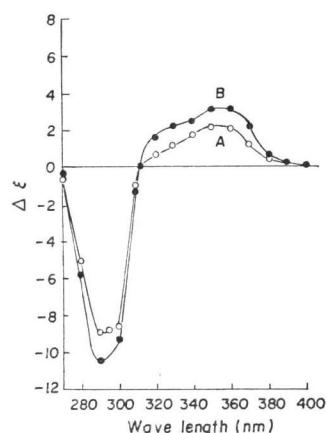


Fig. 2. CD curves of rhodirubins A and B in chloroform - methanol (1: 1)



1000, 980, 960, 920, 800, 790 and 760. The PMR spectrum of rhodirubin B in  $\text{CDCl}_3$  (100 MHz) shows the following chemical shifts (ppm): 7.6, s; 7.24, s; 5.50, m; 5.02, m; 4.84, m; 4.52, q; 4.7 ~ 3.90, overlapping m; 3.72, s; 3.6 ~ 0.09 overlapping m and 2.18, s.

In order to determine the sequence of the hexoses in the sugar chain, rhodirubins A and B were hydrolyzed under three different conditions. On mild hydrolysis with 0.5% HCl at 20°C for 10 minutes both rhodirubins A and B liberated L-rhodosamine. On methanolysis in methanol containing 0.01 N HCl for 30 minutes at room temperature pyrrromycin<sup>3)</sup> (rhodosaminyl- $\epsilon$ -pyrrromycinone) was obtained from both A and B. On total acid hydrolysis (0.1 N HCl for 60 minutes at 85°C)  $\epsilon$ -pyrrromycinone, L-rhodosamine, 2-deoxy-L-fucose and L-rhodosamine were obtained from rhodirubin A and L-rhodosamine was obtained from rhodirubin B in place of 2-

deoxy-L-fucose from rhodirubin A. Thus, the structures shown in Fig. 1 are proposed for rhodirubins A and B. Detailed structural studies will be reported elsewhere.

Among the known anthracycline antibiotics, rhodomycins X and Y<sup>7)</sup>, violamycins<sup>10)</sup> and the compound MA144-N1 have the same sugar moiety (rhodinosyl-2-deoxyfucosyl-rhodosaminyl) as rhodirubin A, but their aglycones were different: rhodomycinones in rhodomycins X and Y and violamycin; aklavinone in MA144-N1;  $\epsilon$ -pyrromycinone in rhodirubins. The sugar moiety (rhodinosyl-rhodosaminyl) of rhodirubin B has not been found in anthracycline antibiotics.

On the stereochemistry of aglycones of anthracycline antibiotics, BROCKMANN *et al.*<sup>8)</sup> have reported the absolute configuration of  $\epsilon$ -pyrromycinone, and ECKARDT *et al.*<sup>9)</sup> have proposed two stereoisomers having the aklavinone structure. The CD curves of rhodirubins A and B are shown in Fig. 2. The same absolute configuration (7S, 9R and 10R) as  $\epsilon$ -pyrromycinone proposed by BROCKMANN *et al.* has been confirmed by the PMR analysis.

The antimicrobial activity of rhodirubins A and B was tested by the broth dilution method with the results shown in Table 3. These antibiotics are strongly active against Gram-positive bacteria. Rhodirubins A and B (2.5~5 mg/kg/day) prolonged the survival period of CDF<sub>1</sub> mice inoculated with L-1210 leukemia. Their LD<sub>50</sub> values in mice were 7.5~10 mg/kg and 10~12.5 mg/kg, i.p., respectively.

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