# COMPOSITION OF THE RIGID-LAYER OF *STREPTOMYCES ANTIBIOTICUS* PRODUCING PEPTIDE ANTIBIOTICS

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Streptomyces antibioticus producing antimycin A and actinomycin did not fit in cell-wall types of aerobic actinomycetes. In addition to universal actinomycetes cell-wall components, *i. e.* glucosamine, muramic acid, glutamic acid and alanine, diaminopimelic acid (DAP), lysine, leucine (isoleucine), glycine, arabinose, glucose, galactose and mannose were detected as major constituents. Phase of the life-cycle influenced the distribution of serine, aspartic acid, lysine and DAP isomers. LL-DAP was found rather than *meso* form. *Meso*-DAP was consistently revealed after 48 hr incubation and not at other times, and as distinct from LL was not strict alternative with lysine. The occurrence of *meso*-DAP, relatively small amounts of murein glucosamine  $(2.2 \sim 5.5 \%)$ and lysozyme resistance suggest a similarity to nocardiae cell-walls. Decrease of murein hexosamine was unfavourable for antimycin A yields. Variation of the murein amino acids stimulated production of actinomycin.

Our understanding of microbial cell-walls has developed rapidly with contributions from a number of laboratories. However, little information is available for groups of organisms other than bacteria. The present paper deals with the nature of cell-walls of *Streptomyces antibioticus* during submerged fermentation of antimycin A and actinomycin. It was felt that the results might provide a sounder basis for the study of comparative biochemistry of streptomycetes and establish some relationships between cell structures and the biosynthesis of secondary metabolites.

### **Materials and Methods**

### Organism and Culture Conditions

The strain of *Streptomyces antibioticus* NRRL 2838 obtained from the Northern Utilization Research and Development Division, Peoria, Ill., was used in experiments. Cultivation on a rotary shaker (240 rpm, 5.3 cm (2.1 in.) amplitude) was carried out at 28°C in flasks (500 ml) with complex medium (150 ml) previously described<sup>9)</sup>.

Preparation of Cell-walls and Rigid-layer (R-layer)

After incubation, the cells were kept in 0.5 % formaldehyde overnight at  $20^{\circ}$ C, collected by centrifugation and washed with 1% HCl and distilled water. After treatment with ethanolic KOH (0.5 %) at 28°C for 24 hours the cells were washed with ethyl alcohol and distilled water. The cells were disrupted by sonication in 0.3 M sucrose solution using an MSE ultrasonic sonicator (20 kc/s) for 6 minutes. The degree of breakage was indicated by the gram stain. The unbroken cells were removed by centrifugation at  $2,000 \times g$ 

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for 20 minutes. The supernatant fluid was then centrifuged at  $17,500 \times g$ , the deposited crude broken cell-walls were washed three times with 0.05 M phosphate buffer pH 7.6 and three times with distilled water. The absence of gross cytoplasmatic materials in cell-walls was checked with the electron microscope (UPT-ČSAV, Brno).

R-Layer of the cell-walls was prepared as follows: The crude cell-walls were resuspended in a freshly prepared and filtered solution of trypsin in 0.05 M phosphate buffer pH 7.6 (0.5 mg/ml) and digested at 37°C for 3 hours on a shaker. Trypsin was removed by washing with phosphate buffer and distilled water. The cell-walls were then subjected to the action of a freshly prepared and filtered solution of pepsin in 0.02 N HCl (1 mg/ml). The mixture was put on the shaker at 37°C overnight, then centrifuged at 17,000 × g for 10 minutes and finally washed with 0.02 N HCl, distilled water and 95 % ethyl alcohol. The purified R-layer preparations were freeze-dried and stored until used.

Analytical Methods

For amino acids analysis the samples were hydrolyzed in 6 N HCl in sealed Pyrex tubes at 105°C for 18 hours. For amino sugars hydrolysis with 7 N HCl at 100°C for 4 hours (FISCHER & NEBEL, 1955) was used. The hydrolysates were filtered, evaporated to dryness three times to remove the HCl and taken up in 0.3 ml of distilled water. Amino acids were detected by electrochromatography<sup>10</sup>, glucosamine and muramic acid by paper chromatography<sup>8</sup>. For quantitative determination of glucosamine ROSENLUND's method<sup>16</sup>) was used.

Diaminopimelic acid (DAP) was identified in the samples hydrolysed with 6 N HCl under reflux for 24 hours<sup>21)</sup>. The washed hydrolysate (15  $\mu$ l equivalent to 2.5 mg dry R-layer) was applied to Whatman No. 1 paper, and one-dimensional descending chromatography was carried out by overnight irrigation with methanol-water-10 N HCl-pyridine (80:17.5:2.5:10, v/v) according to HOARE & WORK<sup>7)</sup>. Chromatograms were dipped in ninhydrin (0.1 % w/v) in acetone and heated for 2 minutes at 100°C (RHULAND *et al.*)<sup>14)</sup>. The DAP spots were light olive green and on standing changed to a stable yellow color with a faint pink fluorescence under ultraviolet light. Since the chromatography did not differentiate between *meso-* and DD-DAP, these isomers were identified enzymatically by means of diaminopimelic acid decarboxylase<sup>7)</sup>. Crude preparations of the enzyme was obtained from *Escherichia coli* NCIB 8113 (CASIDA & BALDWIN)<sup>4)</sup>.

For sugars other than amino sugars samples were hydrolyzed in  $2 \text{ N H}_2\text{SO}_4$  in sealed Pyrex tubes at 100°C for 6 hours. The hydrolysate was neutralized with BaCO<sub>8</sub>. The precipitate of BaSO<sub>4</sub> was centrifuged and the supernatant fluid was evaporated to dryness *in vacuo*. The final product was redissolved in distilled water so as to contain the sugars from 20 mg (dry weight) of the preparation in 0.25 ml. An amount of 10  $\mu$ l of the hydrolysate was spotted on Whatman No. 1 paper for ascending chromatography with 3 successive runs in the solvent system *n*-butanol-acetic acid-water (4:1:5). A standard solution mixture (10  $\mu$ l) of sugars was spotted on the same paper for reference. The papers were dried at room temperature and sprayed with reagent containing 4 % solution of AgNO<sub>8</sub> and 5 N ammonia (1:1).

Susceptibility of the cell-wall material to lysis by lysozyme was determined by suspending the cell-wall preparation in 0.07 M phosphate buffer pH 6.6, containing 0.017 M NaCl, adding egg white lysozyme (100  $\mu$ g/ml) and incubating at 37°C. Decrease in optical density during 2-hour incubation was determined at 20 minutes intervals by the recording spectrophotometer (Unicam SP-700). The antibiotic activity was determined by a microbiological plate method<sup>13</sup> using *Candida albicans*, SZU/44 strain, and *Bacillus subtilis*, FDA 6633 strain, as test organisms and antimycin A and actinomycin C as standards. The representative ultraviolet and visible spectra of partially purified antimycin A and actinomycin<sup>3</sup> were examined in a spectrophotometer (Unicam SP-700).

#### Results

The preparation of the R-layer<sup>20)</sup> of the submerged culture of *S. antibioticus* appeared free from electron dense cytoplasmatic material and has a relatively homogenous appearance without any obvious fine structure (Plate 1). Hexosamine determi-

nation revealed the presence of relatively small amounts<sup>15)</sup> of glucosamine (Fig. 1). Evidence for the 3-O-carboxyethyl derivate of glucosamine, muramic acid, was also shown in the R-layer hydrolysates. Sugar other than aminosugars found in the investigated material included glucose, galactose, mannose and arabinose. No significant difference were found between the individual mycelia of different age in regard to the concentration of reducing sugars except for hexosamine. As shown in Fig. I the apparent decrease in the R-layer hexosamine was reflected as unfavourable for antimycin A yields. Although the presence of glucosamine and muramic acid was a

constant feature of the cell-walls, lysozyme was without action upon *S. antibioticus*. As was the case with whole cell the isolated cellwalls resisted dissolution by lysozyme.

In addition to the common cellwall amino acids, i.e. alanine, glutamic acid, glycine, serine, aspartic acid, lysine and diaminopimelic acid (DAP), leucine (isoleucine) was also found in the S. antibioticus All mentioned amino R-layer. acids were present in a bound form so as to be apparent only after The analytical data hydrolysis. presented in Table 1, Figs. 1 and 2 indicate that at different times of cultivation the R-layer was heterogenous in respect to amino acid composition. Regarding the variation in distribution, attention can be directed to serine, aspartic acid, lysine and DAP isomers (Table l, Fig. 2). The amino acid variation

Plate 1. Electron photomicrograph of the S. antibioticus R-layer, shadowed with chromium at an angle 25 degrees,  $20,000 \times$ 

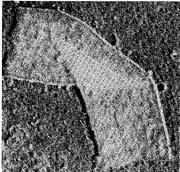


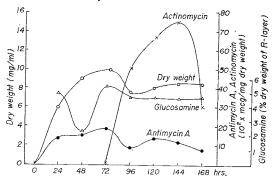
Table 1. Amino acids in R-layer hydrolysates ofS. antibioticus during fermentation ofantimycin A and actinomycin

Time (hrs)	Leu (Ileu)	Ala	Gly	Glu	Ser	Asp	Lys	DAP
24	++	++	++	++	++	++	—	++
48	+	++	+	++	++	+	+	++
72	+	++	+	++	-	+		++
96	+	++	+	++	—	±		++

Note: Except for isoleucine (Ileu) and diaminopimelic acid (DAP) amino acids were referred by the first three letters of their names.

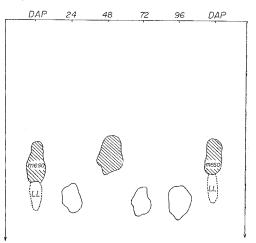
++ an intensely colored spot on chromatogram; + a small definite spot;  $\pm$  doubtful; - no spot, amino acid absent.

Fig. 1. Glucosamine in R-layer of *S. antibioticus* during fermentation of antimycin A and actinomycin



was associated with that phase of the lifecycle at which culture growth has almost ceased, lysis of the cells begun and chromopeptide actinomycin appeared in the culture medium (Fig. 1). Thus it was confirmed that the cell-wall character of young cells was altered as they became old. As to DAP, it is worth emphasizing that analysis of several separate R-layer samples gave consistent results for the content of meso isomer in 48-hour mycelium, while 24-, 72- and 96-hour samples were found to contain DAP in its LL form (Fig. 2). Crude decarboxylase preparation from Escherichia coli completely decarboxylated the DAP of 48-hour mycelium, proving the presence of meso and absence of DD isomer<sup>5)</sup>, and had little action on LL-DAP revealed in the above mentioned

- Fig. 2. Chromatogram of DAP isomers in R-layer of S. antibioticus during fermentation of antimycin A and actinomycin
- Note: DAP denote the reference sample of *meso* diaminopimelic acid with the balance of LL; 24, 48, 72 and 96 denote the age (hr) of the analysed culture.



samples. DAP in its optically inactive configuration did not appear to be a strict alternative with lysine as an amino acid constituent of the R-layer.

#### Discussion

S. antibioticus producing antimycin A and actinomycin yielded cell-walls, the R-layer of which contained glucosamine, muramic acid, glutamic acid, alanine, diaminopimelic acid (DAP), lysine, leucine (isoleucine), glycine, arabinose, glucose, galactose and mannose. It can be concluded that the investigated cell-walls contained mucopeptides (PERKINS & ROGERS<sup>11</sup>), more recently called mureins (WEIDEL & PELZER<sup>19</sup>) or glycoamino-peptides (SALTON<sup>18</sup>), resembling those established in cell-walls of all bacteria so far examined (SALTON<sup>18</sup>). However, S. antibioticus does not fit in any of the four cell-wall types of aerobic actinomycetes considered by BECKER, LECHEVALIER & LECHEVALIER<sup>2</sup>) but resembles a strain of Nocardia, sp. LL-SS 1/5, which is supposed to be a link between the nocardiae and streptomycetes. Relatively small amount of cell-wall glucosamine occurred in S. antibioticus (2.2~5.5%) as compared with that in S. griseus (19.9%), S. fradiae (19.8%) or S. bobiliae (12%). Also lysozyme resistance and occurrence of meso-DAP remind one of Nocardiae (ROMANO & SOHLER<sup>15</sup>; HOARE & WORK<sup>7</sup>).

The findings obtained revealed that the cell-wall character of young cells was altered as they became old. It is reasonable to assume that phase of the life-cycle influenced not only the distribution of hexosamine, serine, aspartic acid and lysine but also the proportion of DAP isomers. The presence of *meso* DAP after 48 hours of incubation and not at other times was surprising. It was possible that LL-DAP might be synthetized first from precursor in the LL configuration and the *meso* isomer was subsequently formed by racemization. One can speculate that the conversion was catalyzed by a specific DAPracemase (ANTIA, HOARE & WORK<sup>1)</sup>) as an important step in lysine biosynthesis. This might explain the fact that *meso*-DAP, as distinct from LL, was not a strict alternative with lysine as amino acid constituent of murein layer.

The apparent decrease in R-layer hexosamine was reflected as unfavourable for antimy-

cin A yields prior to the stationary phase of growth. Taking into account these data, previous results<sup>13)</sup> and the dipeptide character of antimycin A, a correlation might be suggested between biosynthesis-degradation processes of the cell-walls and those involved in the biosynthesis and degradation of antimycin A. The variation of the R-layer amino acids and concomitant production of the peptide actinomycin where culture growth has almost ceased and lysis of the cells begun tempt one to assume the lower specificity of the processes in aging cells and an important difference between the biosynthesis of actinomycin peptide and that of proteins.

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