

STRUCTURE AND ABSOLUTE CONFIGURATION OF THE  
POLYENE MACROLIDE ANTIBIOTIC AMPHOTERICIN B

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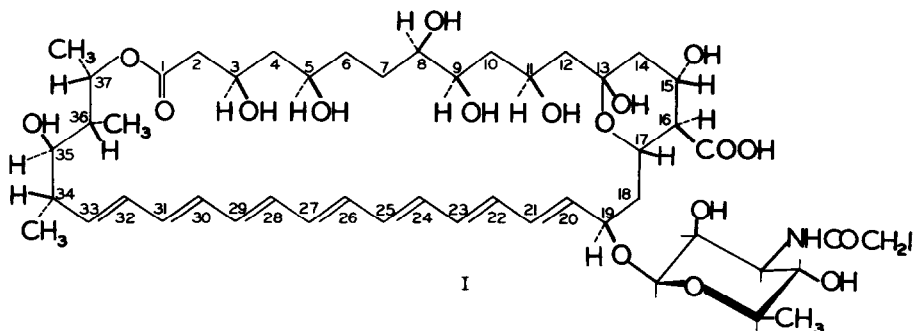
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As a result of the x-ray single crystal analysis of N-iodoacetylamphotericin B the chemical and stereochemical absolute structure is presented (I). A similar structure and stereochemistry was extended to the parent compound itself, amphotericin B, a heptaene macrolide antifungal antibiotic possessing a free amino group in the sugar moiety.



The configuration of the asymmetric centers of the macrolide lactone ring based on the absolute stereochemistry of mycosamine<sup>1,2</sup> are given under the Cahn-Ingold-Prelog system<sup>3</sup> in Table I.

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TABLE

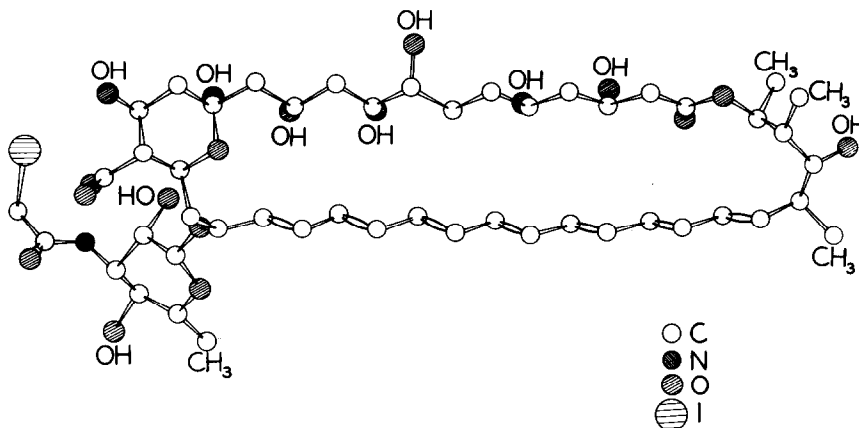
Carbon Atom	3	5	8	9	11	13	15	16	17	19	34	35	36	37
C. I. P. <sup>1</sup> System	R	R	R	R	S	R	S	R	S	R	S	R	R	S

The antifungal antibiotic amphotericin B<sup>4</sup> is produced by Streptomyces nodosus and was first isolated by Vandeputte et al<sup>5</sup> in 1956. Chemical studies by Dutcher<sup>1,2,6</sup>, Borowski<sup>7-11</sup>, Cope<sup>12</sup> and their co-workers, drew out the partial structure of amphotericin B<sup>12</sup>.

Realizing the great difficulties involved in structural studies with the polyene macrolide antibiotics by strictly chemical means we made an attempt to employ the x-ray structure analysis of a single crystal for this purpose. The increasing importance of these antibiotics not only as antifungal agents but also as active substances in the treatment of prostatic hypertrophy<sup>13</sup> and hypercholesterolemia<sup>14</sup> is another reason for determining the total structure of these natural products. The N-iodoacetyl derivative of amphotericin B showing antifungal activity was crystallized from tetrahydrofuran solution. The intensities data were measured with MoK $\alpha$  radiation on a computer controlled automatic Picker diffractometer over a period of 8 days using one crystal. The decrease in standard intensities at the end of the measurements was about 30%. A total of 3702 independent reflections including 2658 non-zero ones were collected within  $2\theta=40^\circ$ .

Crystal data: N-iodoacetylammphotericin B monohydrate tri-tetrahydrofuran, C<sub>49</sub>H<sub>73</sub>O<sub>18</sub>N I H<sub>2</sub>O-3C<sub>4</sub>H<sub>8</sub>O; Mol. wt. ; 1323.9; monoclinic,  $a=21.28$ ,  $b=8.78$ ,  $c=18.69$  Å,  $\beta=103^\circ58'$ ,  $V=3387$  Å<sup>3</sup>,  $D_m=1.33$ ,  $Z=2$ ,  $D_c=1.30$ , Space Group - P2<sub>1</sub>.

After correction for the decrease in intensities, the structure was solved by the heavy atom method and has been refined to an  $R$ -factor of 0.14. The molecular structure of N-iodoacetylammphotericin B as viewed along  $b$ -axis is represented (II). We consider amphotericin B, the parent compound of N-iodoacetylammphotericin B, to be similar in structure to that shown (I). The only difference appears to be the N-iodoacetylation of the free amino group of the antibiotic. In the chemical structure the most interesting feature is the six-membered ketal ring formed from a ketone group at C-13 and the hydroxyl group at C-17.



## II. The molecular structure of N-iodoacetylamphotericin B as viewed along b-axis.

The amino-sugar moiety as a pyranoside is  $\beta$ -glycosidically bound to the hydroxyl at C-19. It is interesting to speculate about the similarity of the amphotericin B moiety between carbon C-13 and C-19 in comparison with the corresponding moieties in the structures of nystatin<sup>15, 16</sup>, pimaricin<sup>17</sup>, lucensomycin<sup>18</sup>, and tetrins A and B<sup>19</sup>. It seems to be possible that all of these other polyene antibiotics could form a six-membered ketal ring. Infrared spectral studies with nystatin and amphotericin B indicate that the free ketone group might not exist. Considering the biogenesis of amphotericin B the proposed total structure is in agreement with the normal polyketide pathway involving acetate and propionate condensation. The only exception involving the positioning of the hydroxyl group at C-8, might readily be explained by an epoxide intermediate mechanism.

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