

The Constitution of the Antibiotic Flambamycin

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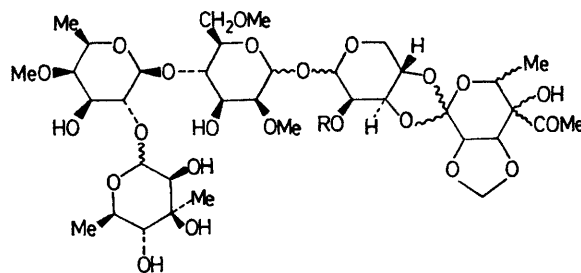
Summary The antibiotic, flambamycin, is shown to have the novel oligosaccharide structure (4) associated with two orthoester linkages.

THE antibiotic, flambamycin, produced¹ by *Streptomyces hygroscopicus* DS 23230 shows high *in vitro* activity against gram-positive bacteria and *Neisseria*. We now report the elucidation of the structure of flambamycin.

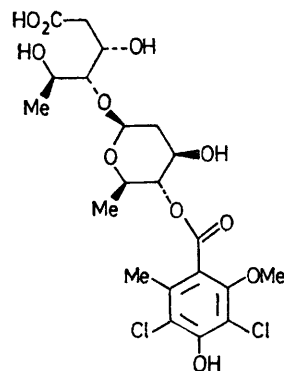
Flambamycin, C₆₁H₈₈Cl₂O₃₃·H₂O, m.p. 202–203 °C (lit.,¹ m.p. 226–228 °C) may be formally regarded as a trianhydro-derivative of isobutyric acid, C₄H₈O₂, flambic acid (1), C₂₁H₂₈Cl₂O₁₁, and flambeurekanose² (2), C₃₈H₅₈O₂₃. Flambamycin forms [acetyltrimethylhydantoin (1.1 mol. equiv.)–methyl cyanide, 12 h, boiling under reflux] a monoacetate, C₆₁H₈₇Cl₂O₃₂(OAc), m.p. 170–172 °C.

The location of the isobutyrate residue as an ester involving position 2 of the L-lyxose residue has been firmly established^{2,3} in flambatriose, flambatetrose, and flambeurekanose isobutyrate. The derivation of the complete structure of flambamycin therefore involves the dehydrative removal of two molecules of water between flambeurekanose monoisobutyrate (3) and flambic acid (1). This, in principle, could be most easily achieved by the occurrence in flambamycin of a second orthoester group: the presence of one orthoester group in flambeurekanose (2) has been firmly established.²

A detailed investigation of the ¹³C n.m.r. spectra of curacin,⁴ flambalactone,³ flambabiose,⁵ flambatriose,⁵ flambatriose isobutyrate,³ flambatetrose,⁵ and flambeurekanose² (2) is highly informative. ¹³C Chemical shift correlations are

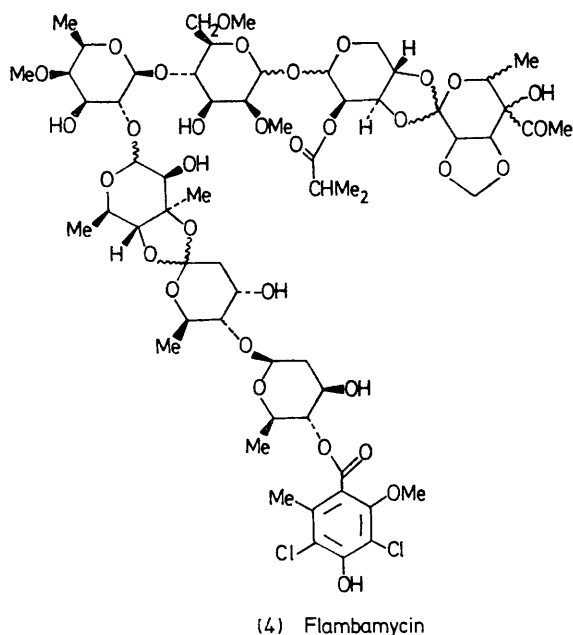


- (2) R = H, Flambeurekanose
(3) R = COCHMe₂, Flambeurekanose isobutyrate



(1) Flambic acid

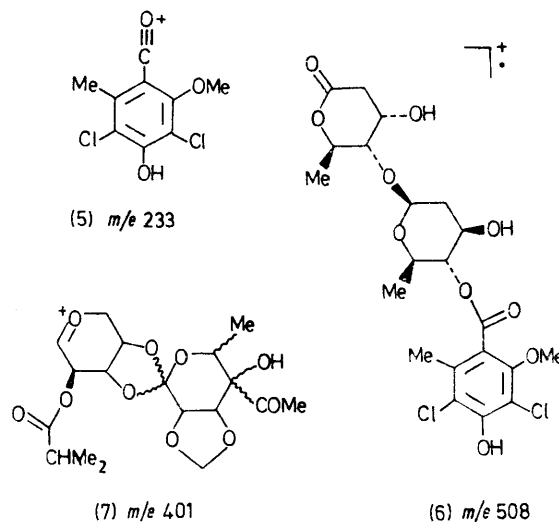
possible between those seven structurally related compounds. In particular, the presence of one orthoester



carbon atom (C_5D_5N , δ 119.8 p.p.m.) in flambeurekanose (2) is clear, whereas in flambamycin, two ^{13}C orthoester signals (C_5D_5N , δ 119.8 and 120.9 p.p.m.) can be assigned. It has been established that the higher relative intensity of the signal (δ 119.8 p.p.m.) in the ^{13}C n.m.r. spectrum of flambamycin is in fact associated with two carbon atoms: one orthoester carbon and one corresponding with an aromatic quaternary carbon atom present either in curacin [(CD_3) $_2CO$, δ 118.9 p.p.m.] or in flambalactone³ (C_5D_5N , δ 119.8 p.p.m.). The second orthoester (δ 120.9 p.p.m.) in flambamycin must be associated with the union between the carboxy-group of flambic acid (1) and flambeurekanose isobutyrate (3). Of the six hydroxy-groups present in flambeurekanose isobutyrate (3), only the three hydroxy-groups associated with the terminal D-evalose residue are sterically suitable for possible involvement in an orthoester linkage. This leads to three possible structures for flambamycin of which the constitution (4) was established as follows.

Permethylation (MeI-NaH-Me₂SO, room temp., 18 h) followed by acidic methanolysis (MeOH-HCl, 3.8% w/v, 1 h, boiling-under reflux) then separation by t.l.c. yielded five new methyl glycosides which were fully characterised

by spectroscopic methods (u.v., i.r., n.m.r.), high-resolution mass spectral fragmentation patterns, specific rotations, and the formation and full characterisation of peroxy-acetates. The degradation products were: (i) curacin O'-methyl ether 3-O-methyl ether methyl glycoside, m.p. 80 °C; (ii) 2-O-methyl-D-evalose methyl glycoside (3,4-di-O-acetate); (iii) 3,4-di-O-methyl-D-fucose methyl glycoside, m.p. 99–101 °C (1,2-di-O-acetate); (iv) 2,3,6-tri-O-methyl-D-mannose methyl glycoside (4-O-acetate); (v) 2-O-methyl-L-lyxose methyl glycoside (3,4-di-O-acetate).



Acidic hydrolysis (5N-HCl, 70 °C, 18 h) of flambamycin (4) gave formaldehyde, isolated from the hydrolysate as its 2,4-dinitrophenylhydrazone (m.p. 164 °C, yield 65%), thus confirming the presence of the methylenedioxy-group. The 1H n.m.r. spectrum of flambamycin was complicated but clearly indicated the presence of four methoxy-groups, one aromatic methyl group, one methyl ketone, one tertiary methyl, and seven secondary methyl groups.

Extensive high-resolution mass spectral studies⁶ were carried out on flambamycin (4) and its derivatives. Major fragment ions (5), (6), and (7) confirmed its structure in important respects, particularly the location of the eurenkanate and isobutyrate residues.

Flambamycin (4) is thus structurally related to the oligosaccharide antibiotics everninomycin-B,⁷ everninomycin-C,⁸ everninomycin-D,⁹ curamycin,⁴ and avilamycin.¹⁰

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