

X-Ray Crystal and Molecular Structure of Antibiotic X-206

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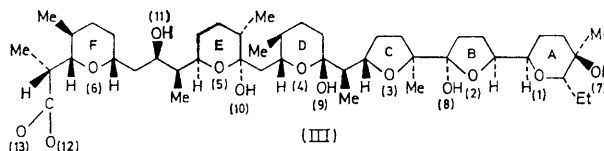
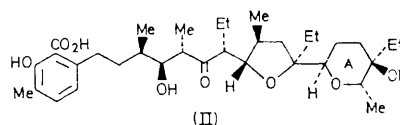
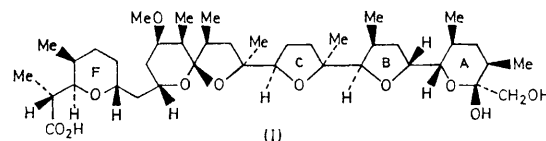
Summary: The molecular structure and absolute configuration of the polyether antibiotic X-206 has been established from an X-ray crystallographic analysis of its silver salt, $C_{45}H_{77}AgO_{13}$.

THE structures of several antibiotics of the polyether type have recently been elucidated by X-ray crystallographic studies of their salts, e.g. monensin¹ and nigericin² (I) which were shown to have similar structures. The isolation of three polyether antibiotics has been reported³ and the first was identical⁴ with nigericin.² The structure⁵ and biosynthesis⁶ of the second (II) were reported recently as was grisorixin.⁷

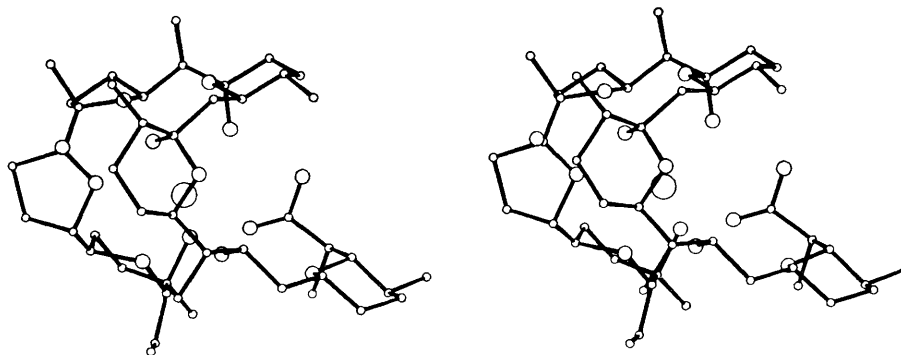
The third (III), antibiotic X-206, ($C_{45}H_{78}O_{13} \cdot H_2O$),⁸ m.p. 133–145°, is a monocarboxylic acid, ν_{\max} (CHCl₃) 1725 cm⁻¹, p*K*_a 8.1 (70% dimethylformamide) and forms a typical organic solvent soluble, water-insoluble sodium salt, m.p. 189–190°, ν_{\max} (CHCl₃) 1570 cm⁻¹ (CO₂⁻).

The structure of X-206 was established as (III) from a three-dimensional X-ray diffraction analysis of its silver salt.³ *Crystal data:* $C_{45}H_{77}AgO_{13}$ $a = 22.90$, $c = 17.44$ Å, $D_m = 1.24$ (floatation in aqueous KI), $Z = 6$, $D_c = 1.19$, space group $P6_3$. 1392 Reflections were measured on a Hilger and Watts Y290 four-circle diffractometer with

established by calculating structure factors for both enantiomers. The results ($R = 8.9$ and 10.2%) show the



molecule to have the configuration depicted in (III) and the Figure.

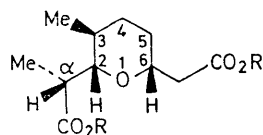


FIGURE

Ni-filtered Cu- K_{α} radiation. Absorption corrections were applied ($\mu = 37$ cm⁻¹). The structure was solved by the heavy-atom method and refined by block-diagonal least-squares. The absolute configuration of the molecule was

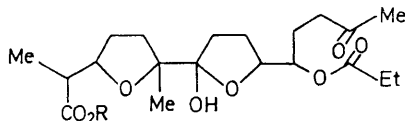
The manner in which the antibiotic ion complexes with the silver ion is similar to that found for other salts of this type.^{1,2,5,7} Here the anion is wrapped around the silver in such a way that its backbone describes a path similar to

that of the seam on a tennis ball. This "seam" is completed by an intramolecular hydrogen bond (2.69 Å) from O(7) to O(13). The silver is co-ordinated unsymmetrically to 6 oxygen atoms [O(1), O(13), O(9), O(11), O(2), and O(10)] with Ag-O distances in the range 2.5–2.8 Å.



(IVa), R = H

(IVb), R = Me



(Va), R = H

(Vb), R = Me

Jones oxidation of (III) gave *inter alia*, the acids (IVa) and (Va). Compound (IVa), m.p. 193°, is an isomer of the dicarboxylic acid isolated^{2b} from nigericin. N.m.r. spectra showed all three ring substituents in (IVa) to be equatorial, δ (CDCl₃) 0.83 (3H, d, *J* 6 Hz, 3-Me), 1.08 (3H, d, *J* 7 Hz, α -Me), 1.00–2.00 (5H, m, $\cdot\text{CH}\cdot\text{CH}_2\cdot\text{CH}_2\cdot$), 2.35 (2H, d, *J*

7 Hz, $\cdot\text{CH}_2\cdot\text{CO}_2$), 2.69 (1H, dq, *J* 3 and 7 Hz, α -H), 3.57 (1H, d, *J* 9.5 and 3 Hz, 2-H), and 3.79 p.p.m. (1H, m, *J* 7 Hz, 6-H). This difference in chemical shift (0.22 p.p.m.) between 2-H and 6-H in both (IVa) and (IVb) is in agreement with the configuration (calc.⁹ 0.28 p.p.m.) given by X-ray rather than the α -C-epimer (calc. 0.94 p.p.m.).

The second oxidation product was isolated as its methyl ester (Vb) after treatment with diazomethane. High resolution mass spectrometry gave a molecular ion at *m/e* 414.2300 consistent with C₂₁H₃₄O₈ and fragment ions at *m/e* 356 (*M* - C₃H₆O) and 355 (*M* - C₂H₂O₂) supporting a methyl ketone and methyl ester respectively.

The two oxidation products were compatible with the structure assigned by X-ray for rings A, B, C, and F. The absolute configuration of (III) shows similarities to the absolute stereochemistry of (II)⁵ and nigericin² (I). After interchanging the methyl and ethyl groups in (III), ring A has the structure identical to the tetrahydropyranyl ring in (II). Three of the four asymmetric centres in ring F [C(2) of (IVa) is the exception] have the same absolute configuration as the analogous centres in nigericin.

Antibiotic X-206 contains twenty asymmetric centres and is the largest polyether antibiotic for which a molecular structure has been established. The molecule is unusual in possessing three lactol functions.

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⁸ Corrected in this paper from C₄₆₋₄₇H₈₀₋₈₂O₁₃ proposed in ref. 3.

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