Vancosamine. A Novel Branched Chain Amino-sugar from the Antibiotic Vancomycin

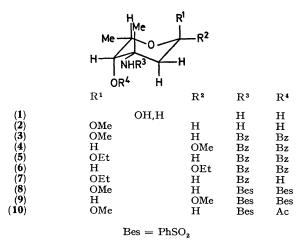
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Summary Acid hydrolysis of the antibiotic vancomycin from Streptomyces orientalis yields the 3-amino-2,3,6trideoxy-3-C-methyl-L-lyxo-hexopyranose, vancosamine, the structure of which is assigned on the basis of chemical and spectroscopic studies, particularly n.m.r. spectroscopy.

WE report the structure and stereochemistry of the branched chain amino-sugar vancosamine (1), $C_7H_{15}NO_3$, which was obtained by acid hydrolysis of vancomycin, an antibiotic derived from *Streptomyces orientalis*.¹ It has previously been shown that acid hydrolysis of vancomycin yielded glucose² and an amino fragment, which was also isolated by acid hydrolyses of the related antibiotics ristomycin, ristocetin, and actinoidin.³ The amine fragment was assigned the impossible formula $C_{12}H_{25}N_2O_5$, and it is probably related to or even identical with, the amino-sugar isolated in the present work.

Hydrolysis of vancomycin with 2N-hydrochloric acid, and separation of the basic compounds using an Amberlite IR-120(H⁺) ion exchange resin yielded a mixture of amines.³



After treatment with methanolic hydrogen chloride, separation of the mixture by t.l.c. $(3\% \text{ NH}_4\text{OH}-\text{Bu}^n\text{OH}$ sat. $\text{H}_2\text{O})$ yielded methyl- α -L-vancosaminide (2). The n.m.r. spectrum ($C_5D_5\text{N}$) contained peaks at τ 7.94 (s, 3-Me) and 8.60 (d, J 6.5 Hz, 5-Me), 5.23 (d, J 4.5 Hz, 1-H), 7.44 (dd, J 4.5, 13.5 Hz, 2ax-H), 7.72 (d, J 13.5 Hz, 2eq-H), 5.97 (s, 4-H), 5.92 (q, J 6.5 Hz, 5-H), and 6.78 (s, OMe), in agreement with the suggested structure. The c.d. spectrum of (2) in 'Cupra A' contained a negative absorption at about 600 nm demonstrating a negative chirality between the *cis*hydroxy- and amino-groups and hence the L-configuration for the amino-sugar.⁴

When the mixture of basic compounds was acetylated using Ac₂O-MeOH, and the product treated with methanolic hydrogen chloride and finally acylated with benzenesulphonyl chloride in pyridine, the acetyl ester (10) was obtained. Thus the initially formed N-acetyl group has migrated to the adjacent *cis*-hydroxy-function under the acidic conditions.⁵

Benzoylation of vancomycin, followed by methanolysis yielded methyl dibenzoyl-α-L-vancosaminide (**3**), m.p. 168— 169°, $\ddagger [\alpha]_D^{3p} - 191°$ (*c* 0·11, MeOH), λ_{max} (MeOH) 227, 270 (infl.) nm (log ϵ 4·44, 3·25), ν_{max} (CHCl₃) 3400, 1725, 1705, 1670 cm⁻¹, τ (CDCl₃) 5·13 (d, *J* 4·5 Hz, 1-H), and a gum, methyl dibenzoyl-β-L-vancosaminide (**4**), $[\alpha]_D^{3p} - 64°$ (*c* 0·14, MeOH), λ_{max} (MeOH) 228, 270 (infl.) nm (log ϵ 4·31, 3·15), ν_{max} (CHCl₃) 3400, 1725, 1705, 1668 cm⁻¹, τ (CDCl₃) 5·36 (dd, *J* 9·5, 2·0 Hz, 1-H).

Irradiation of the C-3 methyl group resonance ($\tau 8.12$) of the dibenzoyl compound (3) caused a 7% NOE (nuclear Overhauser effect), enhancement of the axial C-5 proton signal ($\tau 5.71$, qd, J 6.5, 1 Hz), and no detectable enhancement of the C-4 signal ($\tau 4.89$). Furthermore, on irradiation of the C-3 NH resonance ($\tau 3.37$) there was no observable change in the C-5 proton signal and a small (2—3%) change in the C-4 proton signal, thus confirming the *cis*-diaxial configuration of the C-5 proton and C-3 methyl group.

Benzenesulphonylation of vancomycin followed by methanolysis yielded the corresponding derivatives, methyl

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† C.d. spectra were kindly determined by Dr. P. M. Scopes, Westfield College, London, and by Dr. A. J. McCaffery of this School. ‡ All crystalline compounds had concordant elemental analyses and spectra. dibenzenesulphonyl-α-L-vancosaminide (8), m.p. 132-133°, $[\alpha]_{\rm D}^{24} - 109^{\circ}$ (c 0.34, MeOH) and methyl dibenzenesulphonyl- β -L-vancosaminide (9), m.p. 151–154°, $[\alpha]_{D}^{24}$ – 6.5° (c 0.31, MeOH). Acid hydrolysis of vancomycin followed by evaporation of the crude sugar fraction with added ethanol, and benzoylation of the product yielded ethyl dibenzoyl- α -L-vancosaminide (5), m.p. 131–133°, $[\alpha]_{D}^{25}$ -179° (c 0.27, MeOH) and ethyl dibenzoyl- β -L-vancosaminide (6), m.p. 97–99°, $[\alpha]_{\rm D}^{25} - 82^{\circ}$ (c 0.24, MeOH). The optical rotations of the α -methyl anomers of each of these diacyl derivatives was more negative than the β -

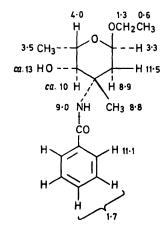


FIGURE. Extrapolated chemical shifts (p.p.m.) for addition of equimolar Eu^{III}(fod)₃ to the benzamide (7).

isomer confirming the assignment of the L-configuration to (1).⁶ The c.d. spectrum^{\dagger} of both (5) and (6) contained negative Davydov bands at 237 nm in agreement with a negative chirality of the two cis-benzoyl groups.7 The positions of the ortho-aromatic proton signals in all the

dibenzoyl compounds were displaced downfield to τ 1.9 indicating that there was an interaction between the cisbenzoyl groups.

Partial alkaline hydrolysis of the dibenzoyl compound (5), τ 4.86 (s, 4-H), yielded the N-benzoyl derivative (7), τ 6.54 (d, J 8.5 Hz, 4-H) and 7.12 (d, J 8.5 Hz, 4-OH, exchanged with D₂O). The n.m.r. spectrum of this monohydroxy-compound was studied using the contact shift reagent $Eu^{III}(fod)_{a}$ and the extrapolated shifts (p.p.m.) for an equimolar solution are shown in the Figure. The lanthanide ion was apparently co-ordinated between the secondary hydroxy-group and the cis-amide oxygen atom.8 The contact shift of the ortho-aromatic signals was particularly marked, whereas the meta- and para-proton signals were barely affected.9

Previous workers had isolated the so-called 'vancomycin acid' Et·CO·CHMe·CH2·CO2H from prolonged acid treatment of vancomycin¹⁰ and in the present work, laevulinic acid has also been isolated. These γ -keto-acids can be regarded as being formed via furans from vancosamine and glucose, respectively.11

Vancosamine can thus be assigned the 3-amino-2,3,6trideoxy-3-C-methyl-L-lyxo-hexopyranose structure (1) and is the first naturally occurring branched chain amino-sugar with geminal methyl and amino groups to be reported. The ¹³C n.m.r. spectrum of (8) fully supports the suggested configuration.12

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Note added in proof: The same structure (1) has been proposed for the amino-sugar from vancomycin by Williams and his co-workers¹³ but without assigning the absolut configuration.

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