

1972, 50, 459-464.

- (5) Insertion products involving monomeric SiF_2 are more common (Liu, C. S.; Hwang, T. L. *J. Am. Chem. Soc.* **1978**, *100*, 2577-2579 and references therein).
- (6) Thompson, J. C.; Wright, A. P. G.; Reynolds, W. F., unpublished work.
- (7) Thompson, J. C.; Margrave, J. L.; Timms, P. L. *J. Chem. Soc., Chem. Commun.* **1966**, 566-567.
- (8) Liu, C. S. Ph.D. Thesis, University of Toronto, Toronto, 1971.
- (9) The observation that the polymer from the *trans*-butene reaction contains the highest proportion of Si_2F_4 units is consistent with the observation that this reaction also gives a higher yield of disilacyclobutane product (III) than any of the other olefin reactions.⁶
- (10) A minor product of empirical formula $\text{C}_2\text{H}_4\text{SiF}_2$ has been detected in the mass spectrum of the products of the reaction of SiF_2 with ethylene.⁷

J. C. Thompson,* A. P. G. Wright, W. F. Reynolds

Department of Chemistry, University of Toronto
Toronto, Ontario, Canada, M5S 1A1

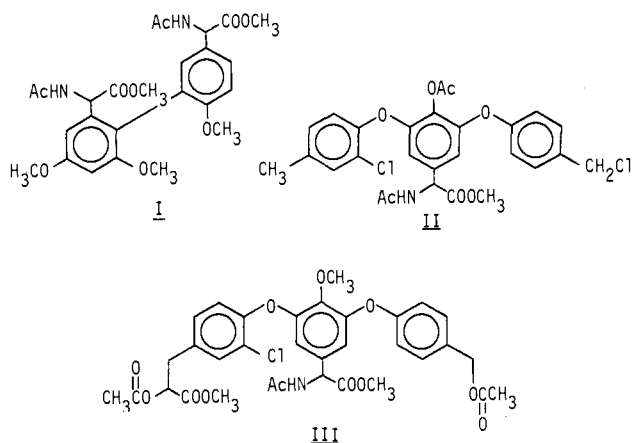
Received December 26, 1978

Avoparcin¹

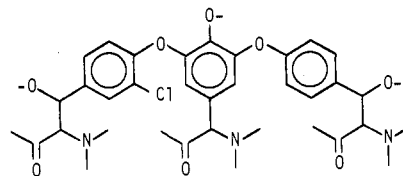
Sir:

Avoparcin, vancomycin, actinoidin, ristomycin, ristocetin, and compound A35512B are a class of complex water-soluble glycopeptide antibiotics with Gram-positive activity which have been isolated over the past 20 years.² Vancomycin is the only member of this antibiotic class the structure of which is known with certainty. Recently Williams at Cambridge prepared modified vancomycin (CDP-I) which was suitable for single-crystal X-ray work and thus obtained the unequivocal structure of this material.³ Avoparcin has commercial importance as a feed additive for agricultural uses. In this communication, we present chemical and spectral evidence that leads us to propose the structure of the aglycone rhamnoside of avoparcin α and β . That avoparcin consists mainly of two components α and β present in about the ratio of 1:3 or 1:4 is readily observed by LC.⁴

Based on the work of Williams et al., we carried out reductive alkaline hydrolysis (refluxing 11 N NaOH, 20% NaBH_4) on avoparcin.⁵ Following suitable derivatization and intensive chromatographic efforts, a number of important fragments were obtained, such as I (M^+ 502, $\text{C}_{25}\text{H}_{30}\text{N}_2\text{O}_9$), II (M^+ 545, $\text{C}_{27}\text{H}_{25}\text{NO}_7\text{Cl}_2$), and III (M^+ 671, $\text{C}_{33}\text{H}_{34}\text{NO}_{12}\text{Cl}$).⁶

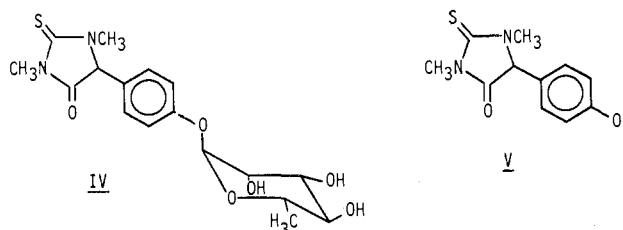


The formation of the aromatic methyl groups as well as the benzyl and lactate moieties in II and III suggests a common origin for these groups. They are reasonably explained in terms of the chemistry of the seryl side chains in the partial structure shown below. Such treatment of avoparcin leads to β elimination of one of the benzylic oxygens which gives rise to an enamide which may be hydrolyzed to a keto acid. Reductive conditions would provide the lactate whereas deoxygenation would lead to the methylbenzene. The benzyl chloride unit

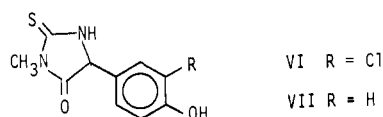


could arise by dealdolization, reduction of the intermediate aldehyde, and introduction of the chlorine during acidification with HCl.

Avoparcin was subjected to the Edman degradation sequence using the reagent methyl isothiocyanate.^{7a,b} At the end of the first stage of the normal Edman two-stage cycle, it was possible to isolate the rhamnoside IV (M^+ 382, $\text{C}_{17}\text{H}_{22}\text{N}_2\text{O}_6\text{S}$).



If the two-stage cycle is completed normally, V (M^+ 236, $\text{C}_{11}\text{H}_{12}\text{O}_2\text{N}_2\text{S}$) is isolated. Subjecting the residue to a second complete Edman cycle yields VI (M^+ 256, $\text{C}_{10}\text{H}_9\text{N}_2\text{O}_2\text{ClS}$) and VII (M^+ 222, $\text{C}_{10}\text{H}_{10}\text{N}_2\text{O}_2\text{S}$) in about the ratio of 3:1 or 4:1. Because of the abnormality at the end of the first stage of



the first Edman cycle,^{7c} these products released at the end of two actual Edman cycles arise because, in effect, three full cycles have been completed. Further the isolation of VI and VII in the ratio mentioned most likely means that the major β component contains the chlorinated *p*-hydroxyphenylglycine compared with *p*-hydroxyphenylglycine in the minor α component.

Although there are obvious differences between the ¹³C NMR spectra of avoparcin (see below) and vancomycin, the general pattern of resonances is similar, especially with respect to the anomeric, aromatic oxycarbon and carbonyl areas, indicating that these antibiotics are structurally related, in agreement with isolation of the identical biphenyl and triphenyl diether (except for chlorine) units from both avoparcin and vancomycin. Further evidence for this relationship comes from the ¹H NMR comparison studies at 270 MHz in $\text{Me}_2\text{SO}-d_6$ (courtesy of Walter Krol, Yale University). An essentially pure sample of the β component and a small sample of mostly the α component were prepared by extensive, repetitive chromatography. Even though the spectra are complex, the two meta-substituted protons on the tetrasubstituted ring of the biphenyl have unique chemical shifts at δ 6.30 and 6.44 in vancomycin and 6.31 and 6.44 in avoparcin α and β .⁸ The α and β curves are almost identical, except in the aromatic region, with the only difference being the extra chlorine in β . Prominent upfield patterns in the α and β spectra are three sharp three-proton doublets at δ 1.11, 1.17, and 1.23. One of these obviously belongs to rhamnose while the other two are assigned to two ristosamine units (see below). The four-proton complex at δ 2.07 is attributed to the C-2 methylene protons of these ristosamines. The *N*-methyl signal of the phenylsarcosine resonates at δ 2.12.

Thus, on the basis of the work described so far, the subunits of the rhamnoside of avoparcin α and β aglycones are depicted,