ACHROMYCIN.¹ THE STRUCTURE OF THE ANTIBIOTIC PUROMYCIN.² I.

Sir:

The new antibiotic, Puromycin, isolated from the mold *Streptomyces alboniger*, has been found to be active against certain bacteria and Trypanosomes.³

active against certain bacteria and Trypanosomes.³ Puromycin, I, m.p. 175.5–177°, $[\alpha]^{25}D - 11^{\circ}$ (ethanol), *Anal.* Calcd. for $C_{22}H_{29}N_7O_5$: C, 56.04; H, 6.20; N, 20.79. Found: C, 56.12; H, 6.48; N, 21.12, is a diacidic base and readily forms a dihydrochloride or a monosulfate. Titration and molecular weight data are in agreement with the above empirical formula. Group analyses show the presence of one amino group (Van Slyke), one methoxyl group, two N-methyl groups and five active hydrogens. A carbonyl group is indicated by a band at 6.05μ in the infrared spectrum. This band may be assigned to a carboxamide grouping. The compound absorbs ultraviolet light with maxima in 0.1 N sodium hydroxide at 275 m μ (E 20,300) and in 0.1 N hydrochloric acid at 267.5 m μ (E 19,500). On acid hydrolysis the ultraviolet absorption maxima are shifted 5 to 10 millimicrons to the longer wave length. Concomitantly, the biological activity of the compound is destroyed.

On alcoholysis with ethanolic hydrogen chloride I is cleaved into three fragments. One of these, II, is an amphoteric compound that precipitates as a dihydrochloride, m.p. 225–227° (dec.), from the cleavage mixture. The free base melts at 257–258°, Anal. Calcd. for C₇H₈N₅: C, 51.52; H, 5.56; N, 42.92; N-methyl, 9.21; mol. wt., Cleavage Tournet, C, 51.56; H, 5.76; N, 43.05; N-methyl, 14.65; mol. wt. (Rast), 169. The analytical data, the ultraviolet and infrared spectra and the amphoteric nature of II suggest a dimethylamino purine. Compound II was identified as 6-dimethylaminopurine by comparison with an authentic sample.

The second fragment, compound III, was identified as O-methyl-L-tyrosine by analysis and by comparison of its melting point, rotation and spectra with those of an authentic sample. The compound was further characterized by representative derivatives.⁵

Compound IV, the third fragment, when isolated as its hydrochloride melts at $158-158.5^{\circ}$ (dec.), $[\alpha]^{25}D - 24.6^{\circ}$ (water), *Anal.* Calcd. for $C_bH_{11}NO_4$ ·HCl: C, 32.35; H, 6.52; N, 7.55; Cl, 19.10; mol. wt., 185.6. Found: C, 32.57; H,

- (1) American Cyanamid Company Trademark for Puromycin.
- (2) Puromycin is the generic name for Achromycin.
- (3) J. N. Porter, R. I. Hewitt, C. W. Hesseltine, G. Krupka, J. A. Lowery, W. S. Wallace, N. Bohonos and J. H. Williams, Antibiotics and Chemotherapy, 2, 409 (1952).
- (4) B. R. Baker, R. E. Schaub, J. P. Joseph and J. H. Williams, to be published.
 - (5) L. Behr and H. T. Clark, This Journal, 54, 1630 (1932).

6.72; N, 7.62; amino nitrogen (Van Slyke), 7.70; Cl, 19.45; neut. equiv., 190.7. The characterizing reactions of IV show a positive Fehling test, a positive Brady's reaction, a negative ninhydrin test and the formation of furfural on deamination and subsequent treatment with phosphoric acid. While IV consumes 3.8 moles of periodic acid in three hours, its N-acetyl derivative reacts with 2.0 moles in the same period. The absence of a carbonyl band in the infrared absorption spectrum of IV (in nujol) and the above chemical and analytical data permit the postulation of IV as a hemiacetal form of a 3- or 4-aminopentose.

The formation of a triacetate of I and its subsequent partial deacetylation with alcoholic ammonia to N-acetylpuromycin permits the postulation of two free alcoholic groups in I. The free amino group in I is placed in the O-methyltyrosine moiety by the failure of I to consume periodic acid. This failure to consume periodic acid by I also eliminates a 4-aminopentose structure for compound IV.

The 3-aminopentose was identified as D-3-aminoribose by comparison of its infrared spectrum, melting point and rotation with a synthetic sample.⁷

A negative Brady's test until after hydrolysis indicates a glycosidic linkage in Puromycin. A comparison of the ultraviolet absorption spectra of I with those of 7- and 9-ethyl-6-dimethylamino-purines⁸ establishes this linkage to be at the 9 position of the purine.

Partial structure V is proposed for Puromycin, I.

Structural features of I to be considered in a future communication are: (1) the α - or β -linkage of the glycoside and (2) the furanosidic or pyranosidic nature of the sugar portion.

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^{(6) 2-}Amino sugars give positive ninhydrin test.

⁽⁷⁾ B. R. Baker, et al., to be published.

⁽⁸⁾ The ultraviolet absorption spectra data for these purines are: 6-dimethylamino-9-ethylpurine $\gamma^{0.1N}_{\max}^{1.N}_{\max}^{1.N}$ 277.5 (E 18,300); $\gamma^{0.1N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\max}^{1.N}_{\min}^{1.N}_{\max}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}_{\min}^{1.N}$