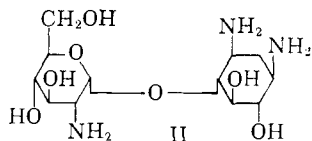


be identical with D-glucosamine hydrochloride by infrared spectrum, X-ray diffraction, optical rotation and comparison of the salicylidene derivatives.

Since the crystalline N,N',N''-triacetylparomamine [Anal. Calcd. for C₁₈H₃₁N₃O₁₀ (449.5): C, 48.10; H, 6.95; N, 9.35. Found: C, 48.04; H, 7.03; N, 9.37; [α]²⁵_D +108° (c 1.0, H₂O); m.p. 300–306° dec.] consumes two moles of periodate⁵ with the absence of formaldehyde formation, and paromamine gave negative reduction tests, the structural formula of the latter is represented as II.



Since the infrared spectra (KBr) of crystalline paromamine trihydrochloride and paromamine free base show absorption bands at 11.2, 11.91 μ and 11.06, 11.75 μ, respectively, no assignment of anomeric configuration is possible by this method.⁶ However, since glucosamine is in the D-series and paromamine trihydrochloride has a high positive molecular rotation (+36,100), an α-D-glycosidic linkage is inferred. Data from methylation experiments described in a subsequent paper have more rigorously established this conclusion.

(5) R. W. Jeanloz and E. Forchielli, *J. Biol. Chem.*, **188**, 361 (1951).

(6) S. A. Barker, E. J. Bourne, M. Stacey and D. H. Whiffen, *J. Chem. Soc.*, 171 (1954); cf. "Methods of Biochemical Analysis," Vol. III, Interscience Publishers, Inc., New York, N. Y., 1956, p. 213.

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RECEIVED MAY 5, 1959

PAROMOMYCIN. II. PAROMOBIOSAMINE, A DIAMINOHEXOSYL-D-RIBOSE

Sir:

In the previous communication¹ methanolysis of the antibiotic paromomycin was reported to yield paromamine and the α- and β-anomers of methyl paromobiosaminide. This communication deals with preliminary structural studies on the disaccharide moiety, paromobiosamine.

The unresolved anomeric mixture of methyl paromobiosaminides was converted to the amorphous N,N'-dibenzoyl derivative [Anal. Calcd. for C₁₁H₁₉N₂O₇(OCH₃)(COC₆H₅)₂: C, 58.64; H, 6.06; N, 5.26. Found: C, 58.37; H, 6.25; N, 5.58] melting over a wide range (120–145°). Dilute acid hydrolysis followed by extraction, ion exchange treatment and carbon chromatography afforded a neutral colorless gum which corresponded to ribose in several paper chromatographic systems. Its infrared absorption spectrum in KBr² was identical to that of D-ribose. Since it exhibited a negative rotation [α]²⁷_D -14° → -18° (18 hr.) the sugar is assigned to the D-series. Reduction

(1) T. H. Haskell, J. C. French and Q. R. Bartz, *THIS JOURNAL*, **81**, 3480 (1959).

(2) F. E. Resnik, L. S. Harrow, J. C. Holmes, M. E. Bill and F. L. Green, *Anal. Chem.*, **29**, 1874 (1957).

with sodium borohydride afforded crystalline ribitol, m.p. 102–103°.

Vigorous acid hydrolysis of methyl paromobiosaminide (6 N hydrochloric acid) resulted in complete destruction of the ribose moiety and, following carbon chromatography, afforded an amorphous hygroscopic diaminoheptose (paromose) [α]²⁶_D +19° (c 1.0, H₂O; no mutarotation). Paromose is characterized as its crystalline dipicrate [Anal. Calcd. for C₆H₁₄N₂O₄ (C₆H₃N₃O₇)₂: C, 33.97; H, 3.17; N, 17.61; picric acid, 72. Found: C, 34.15; H, 3.40; N, 17.44; picric acid (via ultraviolet analysis), 70; [α]²⁵_D +22° (c 0.5, H₂O)] which melted at 126–128° with decomposition. N,N'-Diacetylparomose formed a crystalline p-nitrophenylhydrazone (yellow needles) [Anal. Calcd. for C₁₆H₂₃N₃O₇: C, 48.36; H, 5.83; N, 17.63. Found: C, 48.42; H, 5.82; N, 17.53; [α]²⁵_D +5.9° (c 0.4, moist MeOH), m.p. 229–231° dec.]. N-Acetylation of paromose by the method of Roseman and Ludowieg³ followed by sodium borohydride reduction yielded the crystalline dihydro N,N'-diacetyl derivative [Anal. Calcd. for C₁₀H₂₀N₂O₆: C, 45.45; H, 7.63; N, 10.60. Found: C, 45.29; H, 7.61; N, 10.50; [α]²⁵_D -17.8° (c 4.0, 0.2 M pH 4.5 acetate buffer); m.p. 150.5–151.5°].

Dilute acid hydrolysis (0.5 N hydrochloric acid for 5 hours at 92°) of methyl paromobiosaminide produced, in addition to small amounts of starting material and paromose, the reducing disaccharide paromobiosamine which was isolated as the crystalline dihydrochloride [Anal. Calcd. for C₁₁H₂₂N₂O₆·2HCl·CH₃OH (415.3): C, 34.71; H, 6.80; N, 6.75; Cl, 17.08. Found: C, 34.53; H, 7.13; N, 6.82; Cl, 16.84; neutral equivalent, 204; [α]²⁷_D +25.5° (c 1.0, H₂O; no mutarotation)] and as the free base having a mutarotation value of +32° (c 1.0, H₂O). The ease of methanolysis (0.32 N HCl) of paromomycin to methyl paromobiosaminide which in turn can be hydrolysed with 0.5 N HCl to the free disaccharide, paromobiosamine, argues strongly in favor of a diaminoheptosyl pentose rather than a pentosyl diaminoheptose structure. Rinehart and Woo⁴ arrive at the same conclusion with the neobiosamines but base their evidence on the detection of ribitol from hydrolysis of N,N'-dibenzoyldihydroneobiosamine C.

(3) S. Roseman and J. Ludowieg, *THIS JOURNAL*, **76**, 301 (1954).

(4) K. L. Rinehart, Jr., and P. W. K. Woo, *ibid.*, **80**, 6463 (1958).

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PAROMOMYCIN. III. THE STRUCTURE OF PAROMOBIOSAMINE

Sir:

The characterization of paromobiosamine as an O-(diaminoheptosyl)-D-ribose was described in the previous communication.¹ This report concerns

(1) T. H. Haskell, J. C. French and Q. R. Bartz, *THIS JOURNAL*, **81**, 3481 (1959).