

## Note

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### The sugar sequence of a streptococcal, immunogenic tetraheteroglycan: a revision

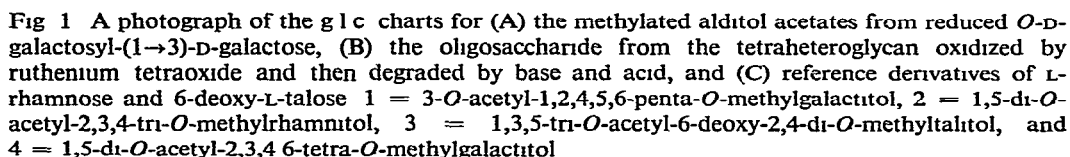
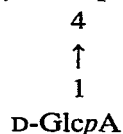
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It was reported recently that an immunogenic tetraheteroglycan from the cell wall of *Streptococcus bovis*, strain C<sub>3</sub>, consists of a main chain of  $\rightarrow 3$ -6-deoxy-L-talose-(1 $\rightarrow$ 3)-L-rhamnose-(1 $\rightarrow$ 3)-D-galactose-(1 $\rightarrow$ 2)-L-rhamnose-(1 $\rightarrow$  and side chains of single D-glucuronic acid groups at O-4 of the 3-substituted rhamnose residues<sup>1</sup>. The immunogenicity of a cell-wall glycan is attributable to the nature of the monosaccharide constituents, the sugar sequence, and the glycosidic bond-types<sup>2</sup>. We have now found that the sequence proposed for the main chain of this glycan needs to be revised to the following  $\rightarrow 3$ -6-deoxy-L-talose-(1 $\rightarrow$ 3)-D-galactose-(1 $\rightarrow$ 3)-L-rhamnose-(1 $\rightarrow$ 2)-L-rhamnose-(1 $\rightarrow$  with side chains of single D-glucuronic acid groups at O-4 of the 3-substituted L-rhamnose residues. This revision involves a reversal of the order of the D-galactose residue and the 3-substituted L-rhamnose residue. It is based on new data from experiments on the chemical degradation of the tetraheteroglycan by alkaline  $\beta$ -elimination, oxidation with ruthenium tetroxide, and degradation by base and acid followed by reduction and methylation analysis of the resulting oligosaccharide.

Fig. 1, frame B, shows a g.l.c. pattern for the methylated alditol acetates from the reduced oligosaccharide. These derivatives were identified as 3-O-acetyl-1,2,4,5,6-penta-O-methylgalactitol, 1,5-di-O-acetyl-2,3,4-tri-O-methylrhamnitol, and 1,3,5-tri-O-acetyl-6-deoxy-2,4-di-O-methyltalitol. The three derivatives possessed retention times identical to those of reference derivatives (frames A and C of Fig. 1), and the derivatives of L-rhamnose and 6-deoxy-L-talose yielded mass-spectral data identical with the data of the reference compounds. The foregoing results establish that the sequence of residues in the oligosaccharide derived from the degraded glycan is L-rhamnose-(1 $\rightarrow$ 3)-6-deoxy-L-talose-(1 $\rightarrow$ 3)-D-galactose. In the native glycan, the D-galactosyl residue is attached to a second L-rhamnosyl residue by a 1 $\rightarrow$ 3 linkage. The second rhamnosyl residue is, in turn, attached by a 1 $\rightarrow$ 2 linkage to the terminal L-rhamnose group of another oligosaccharide and is substituted at O-4 by a D-glucosyluronic acid group. Repetition of this structure several times yields the glycan molecule<sup>1</sup>.

 $\rightarrow 3)-6\text{-deoxy-L-Talp-(1}\rightarrow 3)\text{-D-Galp-(1}\rightarrow 3)\text{-L-Rhap-(1}\rightarrow 2)\text{-L-Rhap-(1}\rightarrow$ 

*Selective oxidation of the tetraheteroglycan with ruthenium tetroxide* — The tetraheteroglycan<sup>3</sup> from *Streptococcus bovis*, strain C<sub>3</sub> (25 mg), was methylated by the Hakomori procedure<sup>4</sup> The methylated glycan was purified by chromatography on Sephadex LH-20 and the purified product subjected to  $\beta$ -elimination to remove the D-glucosyluronic acid groups<sup>5</sup> The product was rechromatographed on Sephadex LH-20, dried in a stream of nitrogen, and oxidized with ruthenium tetroxide<sup>6</sup> A syrupy, methylated glycan devoid of D-glucuronic acid groups was obtained

**Degradation of the methylated and oxidized tetraheteroglycan** — The methylated and oxidized glycan was degraded to an oligosaccharide derivative by treatment first

with base and then with acid, essentially as described in a published procedure<sup>6</sup>. However, in order to effect complete hydrolysis of the product, treatment of the sample with sodium methoxide was extended to 4 h. As revealed in later analyses, some degradation of the reducing-end residue of the oligosaccharide occurred. As a result, methylation analysis gave a lower yield of 3-*O*-acetyl-1,2,4,5,6-penta-*O*-methylgalactitol than of the other derivatives (frame B of Fig 1).

*Structural analysis of the methylated oligosaccharide* — The methylated oligosaccharide, obtained as just described, was reduced to the corresponding alditol with sodium borohydride. The product was dried overnight and then remethylated<sup>4</sup>. The product was successively hydrolyzed, reduced, and acetylated, and the derivatives were identified by g l c -mass spectrometry<sup>3-7</sup>. The results of the g l c analysis are shown in Fig 1, frame B.

*Reference derivatives and analytical methods* — *O*- $\beta$ -D-Galactopyranosyl-(1 $\rightarrow$ 3)-D-galactose<sup>8</sup> (2 mg) was reduced with sodium borohydride, and the product subjected to methylation analysis. The g l c pattern for the derivatives from the reduced oligosaccharide is shown in frame A of Fig 1. L-Rhamnose (5 mg) and the native tetraheteroglycan (3 mg) were methylated and subjected to the conventional series of reactions to obtain methylated alditol acetates<sup>3-7</sup>. The 2,4-dimethyl ether of 6-deoxy-L-talose possessed the shortest retention-time of the derivatives from the tetraheteroglycan. The g l c pattern for this derivative and for the trimethyl ether of L-rhamnose is shown in frame C of Fig 1. The g l c pattern for other derivatives from the tetraheteroglycan is not shown in Fig 1.

G l c analysis was performed in a 1.83 m  $\times$  3.2 mm stainless-steel column packed with 3% OV-225 on 80-100 Supelcoport. A Varian Aerograph 1400 gas chromatograph was used at a temperature of 190°. Mass spectrometry was performed with a Dupont 21-490 mass spectrometer. The source temperature was 240°, the electron potential 70 eV, and the ionization current 125  $\mu$ A. Mass-spectral data were obtained for all derivatives obtained from the oligosaccharide from the glycan except for 3-*O*-acetyl-1,2,4,5,6-penta-*O*-methylgalactitol, which was present in low concentration. The mass-spectral data for the derivatives of L-rhamnose and 6-deoxy-L-talose have been recorded previously<sup>3-7</sup>. Data for the reference compound 3-*O*-acetyl-1,2,4,5,6-penta-*O*-methylgalactitol were as follows (relative abundances in parentheses): *m/e* 45 (100), 89 (80), 101 (90), 133 (20), 205 (15), and 248 (10).

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