

Note

Structure of a glycan from the surface-layer glycoprotein of *Clostridium thermohydrosulfuricum* strain L111-69

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Clostridium thermohydrosulfuricum strain L111-69 and *Clostridium thermosaccharolyticum* strain D120-70 were the first eubacteria for which crystalline surface layers (S-layers)¹ were shown to contain covalently linked carbohydrates². Digestion of the purified *C. thermohydrosulfuricum* glycoprotein with pronase and gel-permeation chromatography³ of the products on Bio-Gel P-30 yielded a single glycopeptide of apparent molecular weight ~20,000.

Carbohydrate analysis by g.l.c. of the alditol acetates after hydrolysis revealed mannose and rhamnose in the molar ratio 1:1.

The ¹H-n.m.r. spectrum of a solution of the glycan in D₂O contained signals for anomeric protons at 4.92 (s, 1 H) and 5.01 p.p.m. (s, 1 H), one signal for a CH₃ group of Rha at 1.27 p.p.m. (d, 3 H), and signals (10 H) in the range 3.4-4.2 p.p.m.

The proton-decoupled ¹³C-n.m.r. spectrum of the glycan contained twelve signals corresponding to twelve carbons (Table I). Thus, the glycan was a regular polysaccharide with a disaccharide repeating-unit. There was one signal at 17.58 p.p.m. corresponding to C-6 of Rha, one at 61.34 p.p.m. (C-6 of Man), eight in the range 67.4-76.4 p.p.m., and two in the anomeric region (97.05 and 101.68 p.p.m.).

The proton-coupled ¹³C-n.m.r. spectrum exhibited sufficient features to establish a hypothetical structure. The *J* value (172 Hz) of each of the C-1 signals

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TABLE I

^{13}C -CHEMICAL SHIFTS^a AND $^1J_{\text{C,H}}$ VALUES (Hz) OF THE *Clostridium thermohydrosulfuricum* L111-69 GLYCAN AND SELECTED MODEL COMPOUNDS

Glycosyl residue	C-1	C-2	C-3	C-4	C-5	C-6
(1→3)- α -Rhap-(1→4)	101.68	67.35	75.25	71.13	70.04	17.58
	172	148	142	144	(144)	127
α -Rhap ^b	94.70	71.54	70.70	72.91	68.99	17.59
	170	149	142	143	145	128
(1→4)- α -Manp-(1→3)	97.05	71.53	70.08	76.44	72.39	61.34
	172	150	(144)	148	(142)	144
α -Manp ^b	94.66	71.46	71.02	67.68	73.20	61.81
	170	149	145	146	142	144

^aIn p.p.m. downfield from the signal of Me₄Si (67.40 p.p.m. upfield from the signal of 1,4-dioxane in D₂O). Couplings were measured by hand on enlarged 62.9-MHz spectra, and those from overlapping signals shown in brackets were determined from the doubled difference between the centre of the signal and one part each of the coupled signals. ^bAssignment from ref. 4.

at 101.68 and 97.05 p.p.m. (Table I) indicated⁵ the pyranosyl residue to be α . Since $^1J_{\text{C,H}}$ values for monosaccharides are not significantly different from those in the corresponding glycosyl units of oligo-⁶ or poly-saccharides⁷, three resonances were selected [67.35 (J 148 Hz), 71.53 (J 150 Hz), and 76.44 p.p.m. (J 148 Hz)] as candidates for C-2 of Rha and Man. However, the first two lacked large two- or three-bond C-H couplings, a feature which is diagnostic of the C-2 resonances of Rha or Man. From the chemical shifts, glycosylation at either C-2 was excluded. Since no additional signal upfield of 70 p.p.m. was observed (unsubstituted C-4 of Man), Man is glycosylated in position 4. Because of the relatively large $^1J_{\text{C,H}}$ value (similar to that of the monosaccharide), the signal at 76.44 p.p.m. (J 148 Hz) is assigned to C-4 of Man.

Considering that C-6 (61.34 p.p.m.) of Man is unsubstituted, C-4 (76.44 p.p.m.) is substituted, both C-2's (67.35 and 71.53 p.p.m.) are unsubstituted, and one C-2 (67.35 p.p.m.) is adjacent to a glycosidic linkage, the following two structures are possible: (a) a (1→4)-mannan having rhamnosyl branches at positions 3, or (b) \rightarrow 4)- α -Manp-(1→3)- α -Rhap-(1→ repeating-units (**1**).

Structure (a) was ruled out by a Smith-degradation experiment since only the Rha units remained intact and low-molecular-weight fragments were obtained.

Next, the C-1 signals (101.68 and 97.05 p.p.m.) were assigned. No large up-field shift is to be expected⁸ for an $\alpha\text{D}_{2a}\rightarrow$ 4D_{2a} ($\alpha\text{L}_{2a}\rightarrow$ 4L_{2a})-linkage⁹ or an $\alpha\text{L}_{2a}\rightarrow$ 4D_{2a}

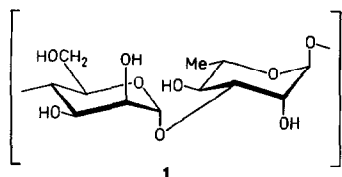


TABLE II

¹H-CHEMICAL SHIFTS^a OF THE *Clostridium thermohydrosulfuricum* L111-69 GLYCAN DETERMINED WITH CHORTLE¹³

Glycosyl residue	H-1	H-2	H-3	H-4	H-5	H-6
(1→3)- α -Rhap-(1→4)	4.92	4.19	3.8	3.51	4.03	1.27
(1→4)- α -Manp-(1→3)	5.01	3.9	3.9	3.8	3.9	3.7 ^b

^aShifts are in p.p.m. downfield from the signal of sodium 3-(trimethylsilyl)propionate-*d*₄. Shifts determined from the proton spectrum directly are given with two decimals. ^bCHORTLE affords only the mean of the chemical shifts of both the H-6 resonances. These signals are located at ~3.45 and ~3.95 p.p.m.

(α D_{2a}→4L_{2a})-linkage¹⁰. Thus, the signal at 101.68 p.p.m. corresponds to C-1 of Rha. On the other hand, for an α D_{2a}→3L_{2a}-linkage¹¹, an upfield shift is to be expected in contrast to an α L_{2a}→3L_{2a}-linkage¹². Therefore, the signal at 97.05 p.p.m. corresponds to C-1 of Man. The remaining signals were assigned by application of the reported substitution shifts¹¹.

The ¹H-n.m.r. spectrum was assigned using the CHORTLE (carbon-hydrogen correlations from one-dimensional polarization-transfer spectra by least-squares analysis)¹³ method (Table II). The observed 0.4-p.p.m. downfield shift of the Rha H-5 resonance agrees with the structure, because H-5 of Rha is close to the HO-3 of Man in the glycan. Similar effects are observed if H-1 is close to an oxygen¹⁴. The same deshielding (0.4 p.p.m.) of H-5 of Man (compared to methyl α -D-mannopyranoside) is observed, because H-5 is close to HO-4 of the Rha.

The [α]_D value of the polysaccharide based on those for monosaccharides was calculated according to Klyne's rule¹⁵ (Table III). The observed [α]_D²⁰ value of +12° (c 1, water) agrees well with the value (+11°) calculated for the D-Manp/L-Rhap combination.

TABLE III

[M]_D VALUES CALCULATED ACCORDING TO KLYNE'S RULE¹⁵ FOR POLYSACCHARIDES COMPOSED OF Man-Rha DISACCHARIDE REPEATING-UNITS

Compound	[α] _D ²⁰ (degrees)	[M] _D (degrees)
Methyl α -D-mannopyranoside ¹⁶	+79.2	+154
Methyl α -L-mannopyranoside ¹⁶	-79.4	-154
Methyl α -D-rhamnopyranoside	+67.2	+120
Methyl α -L-rhamnopyranoside ¹⁷	-67.2	-120
Polysaccharides (calc.)		
(D-Manp + D-Rhap)	+89	+274
(D-Manp + L-Rhap)	+11	+34
(L-Manp + D-Rhap)	-11	-34
(L-Manp + L-Rhap)	-89	-274
<i>C. thermohydrosulfuricum</i> L111-69 glycan, observed	+12.3	+38

EXPERIMENTAL

General. — Hexose and rhamnose were determined colorimetrically or by g.l.c.³ The $[\alpha]_D^{20}$ value was determined with a Perkin-Elmer model 141 polarimeter. ¹H- and ¹³C-n.m.r. spectra were recorded³ with a Bruker WM 250 instrument at 298 K for solutions in D₂O. The CHORTLE experiment¹³ was performed using the pulse sequence described at 298 K for a solution of the glycan in D₂O.

Isolation of the glycopeptide. — *Clostridium thermohydrosulfuricum* strain L111-69 was grown² under anaerobic conditions in TS-medium (Becton Dickinson) in 4-L bottles at 65°. The isolation of the S-layer glycoprotein and degradation into the glycopeptide with pronase followed described methods³. During the purification, chromatography on Dowex 50W-X8 (H⁺) resin was omitted. After chromatography on a column (1 × 113 cm) of Bio-Gel P-4, the final purification was performed on a calibrated column (1 × 113 cm) of Bio-Gel P-30 equilibrated with aqueous 1% acetic acid. Appropriate fractions were collected and lyophilised, and the glycopeptide was stored at -20°.

*Smith degradation*³. — The glycopeptide (18 mg) was oxidised for 48 h at 4° in the dark, then reduced with sodium borohydride, hydrolysed with aqueous 5% acetic acid (1 mL) for 1.5 h at 100°, and applied to a column (1 × 113 cm) of Bio-Gel P-2 equilibrated with water. Fractions (1 mL) were tested³ for rhamnose.

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