

Note

Somatic antigens of pseudomonads: structure of the O-specific polysaccharide of *Pseudomonas syringae* pv. *tomato* 140(R)

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This work is a continuation of a chemical and immunochemical study of lipopolysaccharides of the phytopathogenic bacteria *Pseudomonas syringae* and related species (refs. 1–3 and refs. therein). We now report the structure of the O-specific polysaccharide of *P. syringae* pv. *tomato* 140(R).

The O-specific polysaccharide was obtained by mild acid degradation of the lipopolysaccharide, isolated by washing of bacterial cells with saline⁴. It had $[\alpha]_D +17^\circ$ (*c* 0.5).

Acid hydrolysis of the polysaccharide revealed the presence of rhamnose and 3-amino-3,6-dideoxygalactose (Fuc3N), which were identified by using GLC of alditol acetates and an amino acid analyzer, respectively. GLC of (+)-2-octyl glycosides of rhamnose, as compared with (+)- and (–)-2-octyl glycosides of the authentic sample of L-rhamnose, proved⁵ that this sugar has the L configuration. The D configuration of Fuc3N was established on the basis of NOE and ¹³C chemical shift data (see below).

The ¹³C NMR spectrum of the polysaccharide (Fig. 1, Table I) contained the signals for four anomeric carbons at 97.5–103.2 ppm, one carbon bearing nitrogen at 52.6 ppm, four CH₃–C groups (C-6 of 6-deoxyhexoses) at 16.5 (one) and 18.0–18.2 ppm, other sugar carbons in the region 67.2–79.2 ppm, and one N-acetyl group (Me at 23.4 ppm, CO at 175.6 ppm). In the spectrum, there were no lines characteristic of furanose rings and, hence, the constituent sugars were pyranoid.

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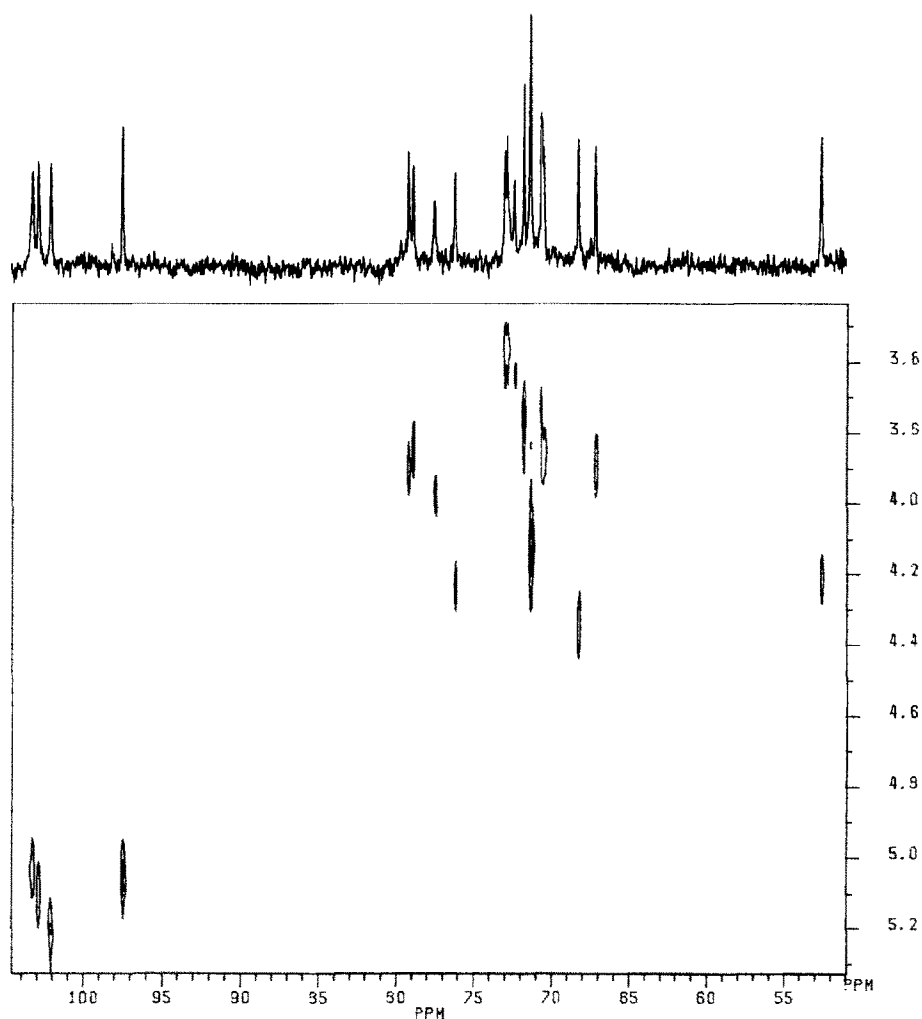


Fig. 1. 75-MHz 2D heteronuclear ^{13}C - ^1H shift-correlated spectrum of the O-specific polysaccharide. The respective part of a 1D ^{13}C NMR spectrum is displayed along the F_2 axis.

TABLE I

^{13}C NMR data for the O-specific polysaccharide (δ in ppm)^a

Unit	C-1	C-2	C-3	C-4	C-5	C-6
$\rightarrow 3)\text{-}\alpha\text{-L-Rhap-(1} \rightarrow$ (A)	102.9	71.4	78.9	73.0 ^b	70.7	18.0 ^d
$\rightarrow 2)\text{-}\alpha\text{-L-Rhap-(1} \rightarrow$ (B)	102.0	76.2	77.5	72.4	70.6 ^c	18.2 ^d
3 ↑						
$\rightarrow 3)\text{-}\alpha\text{-L-Rhap-(1} \rightarrow$ (C)	103.2	71.4	79.2	72.9 ^b	70.5 ^c	18.0 ^d
$\alpha\text{-D-Fucp3NAc-(1} \rightarrow$ (D)	97.5	67.2	52.6	71.8	68.3	16.5

^a Additional signals: NAc at 23.4 ppm (Me) and 175.6 ppm (CO). ^{b,c,d} Assignments could be interchanged.

TABLE II

¹H NMR data for the O-specific polysaccharide (δ in ppm, J in Hz) ^a

H-1	H-2	H-3	H-4	H-5	H-6
$\rightarrow 3)\text{-}\alpha\text{-L-Rhap-(1}\rightarrow$ (unit A)					
5.10	4.10	3.83	3.53	3.73	1.24
$J_{1,2} < 2$	$J_{2,3} \sim 3$	$J_{3,4} \sim 10$	$J_{4,5} \sim 10$	$J_{5,6} 6.5$	
$\rightarrow 2)\text{-}\alpha\text{-L-Rhap-(1}\rightarrow$ (unit B)					
5.19	4.22	3.96	3.61	3.84	1.32
$J_{1,2} < 2$	$J_{2,3} 2.5$	$J_{3,4} 9.6$	$J_{4,5} \sim 10$	$J_{5,6} 6.1$	
$\rightarrow 3)\text{-}\alpha\text{-L-Rhap-(1}\rightarrow$ (unit C)					
5.01	4.13	3.89	3.55	3.84	1.26
$J_{1,2} < 2$	$J_{2,3} \sim 3$	$J_{3,4} 9.5$	$J_{4,5} \sim 10$	$J_{5,6} 5.9$	
$\alpha\text{-D-FucpNAc-(1}\rightarrow$ (unit D)					
5.04	3.88	4.20	3.74	4.33	1.15
$J_{1,2} 4.0$	$J_{2,3} \sim 11$	$J_{3,4} 3.1$	$J_{4,5} < 1$	$J_{5,6} 6.6$	

^a Additional signal: NAc at 2.02 ppm.

The ¹H NMR spectrum of the polysaccharide (Fig. 2, Table II) contained the signals for four anomeric protons at 5.0–5.2 ppm, one of which represented a doublet with $J_{1,2}$ 4.0 Hz and belonged, thus, to H-1 of the α -linked Fuc3NAc. Three other H-1 signals were broadened singlets, and, based on their chemical shifts, one could conclude that all Rha residues are also α -linked. Four CH₃–C groups resonated at 1.1–1.35 ppm (doublets with $J_{1,2}$ 5.9–6.5 Hz), other sugar protons in the region 3.4–4.1 ppm, and the *N*-acetyl group at 2.02 ppm (s).

These data showed that the polysaccharide has a tetrasaccharide repeating unit, which includes three residues of rhamnose and one Fuc3NAc residue, all of them being in the α -pyranoid form.

The ¹H NMR spectrum of the polysaccharide was assigned with the use of selective spin-decoupling procedures, 2D homonuclear shift-correlated (COSY) and one-step relayed coherence transfer (COSYRCT) spectroscopy (Table II). As expected, the doublet at 5.04 ppm was identified as belonging to H-1 of the α -linked Fuc3NAc residue and three broadened singlets to H-1 of the rhamnose residues.

Pre-irradiation of H-1 of one of the rhamnose residues (unit A) at 5.10 ppm resulted in a considerable NOE response on H-2 of another rhamnose residue (unit B) at 4.22 ppm. In turn, in the spectrum obtained with pre-irradiation of H-1 of unit B at 5.19 ppm, there were observed an NOE on H-3 of the third rhamnose residue (unit C) at 3.89 ppm, and pre-irradiation of H-1 of unit C at 5.01 ppm caused an NOE on H-3 of unit A at 3.83 ppm (Fig. 2). These data showed that three rhamnose residues form the main chain of the polysaccharide, where units A and C are substituted at position 3 and unit B at position 2.

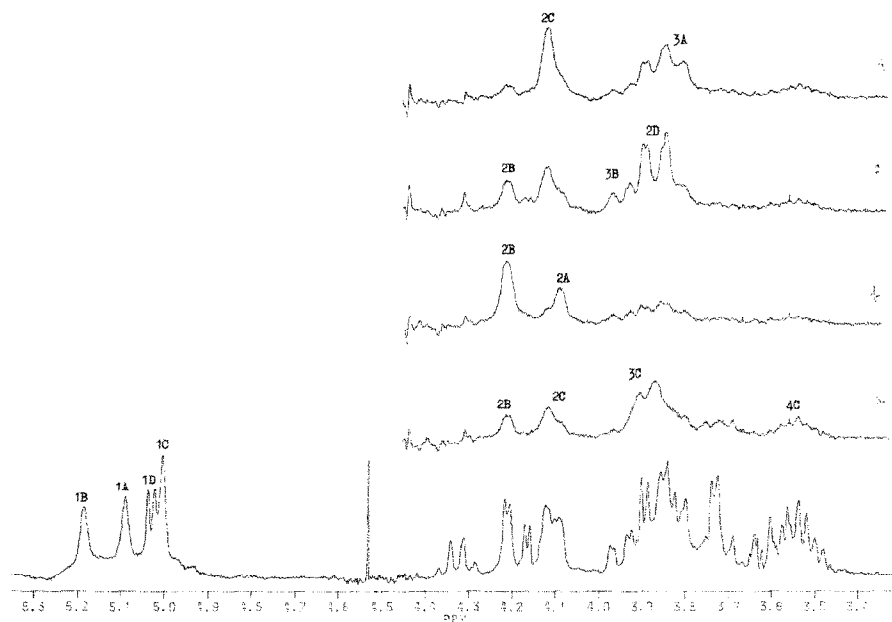


Fig. 2. Part of a 250-MHz ^1H NMR spectrum of the O-specific polysaccharide (bottom curve) and NOE spectra with pre-irradiation of H-1 of units **B**, **A**, **D**, and **C** (curves a, b, c, and d, respectively).

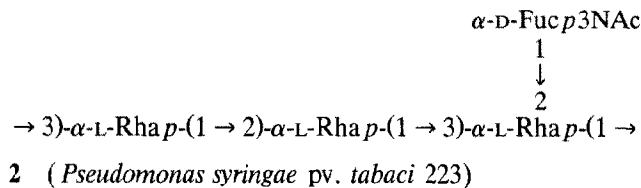
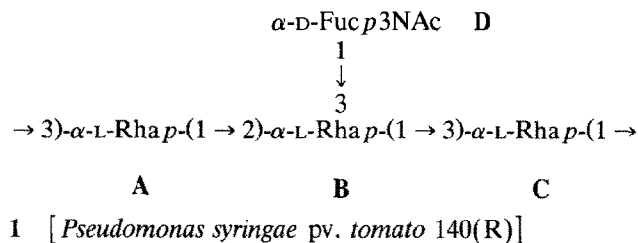
Besides the above-mentioned responses, in the NOE spectra, there were observed considerable responses on H-2 (but not H-3 and H-5) of each of the pre-irradiated sugar residues, which confirmed the α configuration of the rhamnopyranosyl linkages. Only small NOEs, which were caused by spin diffusion characteristic of polymers, arose on H-2 of units **A** and **C** as a result of pre-irradiation of H-1 of units **C** and **B**, respectively. This was additional evidence of the same (L) absolute configuration of the three rhamnose residues since, in the α -(1 \rightarrow 3)-linked disaccharide of rhamnopyranoses with different absolute configurations, the NOE on H-2 is known⁶ to be twice as intense as that on H-3.

Furthermore, pre-irradiation of H-1 of the FucNAc residue (unit **D**) caused considerable NOEs on H-2 and H-3 of unit **B**, proving the attachment of unit **D** as a side chain to unit **B** at position 3. The appearance of the NOE on both H-2 and H-3 is typical⁶ of disaccharide units with L-rhamnopyranose, 3-substituted by an α -pyranoside having the opposite absolute configuration (i.e., the D configuration).

Assignment of the ^{13}C NMR spectrum of the polysaccharide, by using 2D heteronuclear ^{13}C – ^1H shift-correlated spectroscopy (Fig. 1, Table I), was in agreement with the results of the NOE linkage analysis. In particular, the relatively low-field positions of the signals for C-3 of units **A** and **C** at 78.9 and 79.2 ppm, respectively, and for C-2 and C-3 of unit **B** at 76.2 and 77.5 ppm, respectively, as compared to their positions in the spectra of the corresponding free monosaccharides, were due to the α -effects of glycosylation⁷ and confirmed independently the substitution patterns of the sugar residues established by using NOE spectroscopy.

(see above). The high-field position of the signal for C-1 of the Fuc3NAc residue at 97.5 ppm, which corresponded to a small glycosylation effect (ca. 4 ppm), confirmed^{8,9} the different absolute configurations of Fuc3NAc and rhamnose, confirming the D configuration of Fuc3NAc (a chemical shift of ~ 101 ppm would be expected if Fuc3NAc and rhamnose had the same absolute configuration).

The data obtained showed that the O-specific polysaccharide of *P. syringae* pv. *tomato* 140(R) has the structure **1**.



The O-antigen of pv. *tomato* clearly belongs to the structural type characteristic of O-antigens of other *P. syringae* pathovars studied previously (refs. 1–3 and refs. therein), which consist of a linear rhamnan or have a rhamnan backbone with a monosaccharide side chain. It is most closely related to the O-antigen of pv. *tabaci* (structure **2**), which has the same backbone and lateral Fuc3NAc residue but differs in the site of attachment of the side chain¹⁰. Such a similarity of the O-antigens substantiates serological cross-reactivity of these two strains, which served as a basis of their placing¹¹ in the same serogroup VII.

O-antigens of *P. syringae* pathovars belonging to serogroup I possess the same Fuc3NAc side chain but differ in the structure of the rhamnan main chain, which lacks strict regularity, being built up of tetrasaccharide units of two types^{9,12}. The O-antigens of various pathovars belonging to serogroup I differ from each other in the same manner as those belonging to serogroup VII. It is noteworthy that the backbone of the O-antigens of pvs. *tomato* and *tabaci* (structures **1** and **2**) is identical to that of some pathovars of serogroup II (refs. 1,13,14), except that it contains L-rhamnose instead of D-rhamnose.

EXPERIMENTAL

Optical rotation was measured with a Jasco DIP 360 polarimeter for a solution in water at 25°C.

^1H NMR spectra were recorded with a Bruker WM-250 instrument for solutions in D_2O at 55°C . ^{13}C NMR spectra were recorded with a Bruker AM-300 instrument for solutions in D_2O at 60°C . Acetone was used as an internal standard (δ_{H} 2.23 ppm, δ_{C} 31.45 ppm). Selective spin decoupling, 1D NOE, and 2D homonuclear ^1H – ^1H and heteronuclear ^{13}C – ^1H shift-correlated spectroscopy (COSY) were carried out as described^{15,16}.

Growth of bacteria, isolation of lipopolysaccharide, and isolation of O-specific polysaccharide were performed as described^{4,17}.

For sugar analysis, the polysaccharide was hydrolysed with 2 M trifluoroacetic acid (100°C , 2 h), and the hydrolysate was analysed by using GLC of alditol acetates and an amino acid analyzer⁹.

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