



# Structure of the LPS O-chain from *Fusobacterium nucleatum* strain ATCC 23726 containing a novel 5,7-diamino-3,5,7,9-tetradeoxy-L-glucosonic acid presumably having the D-glycero-L-glucosonic configuration

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## ABSTRACT

*Fusobacterium nucleatum* is an anaerobic bacterium found in the human mouth where it causes periodontitis. It was also found in colorectal cancer tissues and is linked with pregnancy complications, including pre-term and still births. Cell surface structures of the bacterium could be implicated in pathogenesis. Here we report the following structure of the lipopolysaccharide O-chain of a spontaneous streptomycin resistant (SmR) mutant of *F. nucleatum* strain ATCC 23726:

-4-β-Non5Am7Ac-4-β-D-GlcNAcyl3NFoAN-3-β-D-FucNAc4N-

where GlcNAcyl3NFoAN indicates 2,3-diamino-2,3-dideoxyglucuronic acid amide with Fo at N-3 being formyl and Acyl at N-2 being propanoyl (~70%) or butanoyl (~30%); Non5Am7Ac indicates 7-acetamido-5-acetimidoylamino-3,5,7,9-tetradeoxy-L-glucosonic acid presumably having the D-glycero-L-glucosonic configuration. To our knowledge, no L-glucosonic isomer of higher sugars of this class as well as no N-propanoyl or N-butanoyl group have so far been found in bacterial polysaccharides.

## 1. Introduction

*Fusobacterium nucleatum* is a Gram-negative anaerobic bacterium found in the human mouth where it can cause periodontitis. It was also found in colorectal cancer tissues, where it promotes chemoresistance [1] and is linked with pregnancy complications, including pre-term and still births [2,3]. *F. nucleatum* has also been implicated in a wide variety of systemic diseases, including GI disorders [4], atherosclerosis [5] and respiratory tract infections [6].

Bacterial surface polysaccharides can be involved in pathogenesis, interacting with immune system. For this reason we have been systematically analyzing *F. nucleatum* LPS O-chain polysaccharides (OPS). Several structures have been determined, showing the presence of unusual monosaccharides and unusual acyl groups [7–10]. Here we report structure of the *F. nucleatum* strain ATCC 23726 SmR OPS, which contains a novel isomer of 5,7-diamino-3,5,7,9-tetradeoxynon-2-ulonic acid.

## 2. Results and discussion

Lipopolysaccharide was extracted from *F. nucleatum* strain ATCC 23726 SmR with hot aqueous phenol and hydrolyzed with 2% HOAc to give an OPS, a LPS core with one OPS repeating unit (core-RU) and other fractions isolated by gel permeation chromatography on a Biogel P6 column. Monosaccharide analysis of the OPS by GC-MS of the alditol acetates showed the presence of the components of the core (glucose, galactose, L-glycero-D-manno-heptose, glucosamine).

2D NMR spectra (<sup>1</sup>H–<sup>1</sup>H gCOSY, <sup>1</sup>H–<sup>1</sup>H TOCSY, <sup>1</sup>H–<sup>1</sup>H NOESY, <sup>1</sup>H–<sup>13</sup>C HSQC (Fig. 1), <sup>1</sup>H–<sup>13</sup>C HMBC) of the OPS contained signals of three sugar spin systems of 2,3-diamino-2,3-dideoxyglucuronic acid (GlcN3NA, A), 2,4-diamino-2,4,6-trideoxygalactose (FucN4N, B), and 5,7-diamino-3,5,7,9-tetradeoxy-L-glucosonic acid (Non, X). The core contained phosphocholine, and a strong signal of its methyl groups was present in all spectra. Configurations of the monosaccharides A and B were inferred from a TOCSY signal pattern and <sup>1</sup>H and <sup>13</sup>C NMR signal positions (Table 1). For the determination of the configuration of Non, better resolved NMR spectra of the core-RU were used, in which this sugar occupied the non-reducing terminal position.

**Abbreviations:** Am, acetimidoyl; Fo, formyl; FucNAc4N, 2-acetamido-2,4,6-trideoxy-galactose; GlcN3NA, 2,3-diamino-2,3-dideoxyglucuronic acid; Non, 5,7-diamino-3,5,7,9-tetradeoxy-L-glucosonic acid

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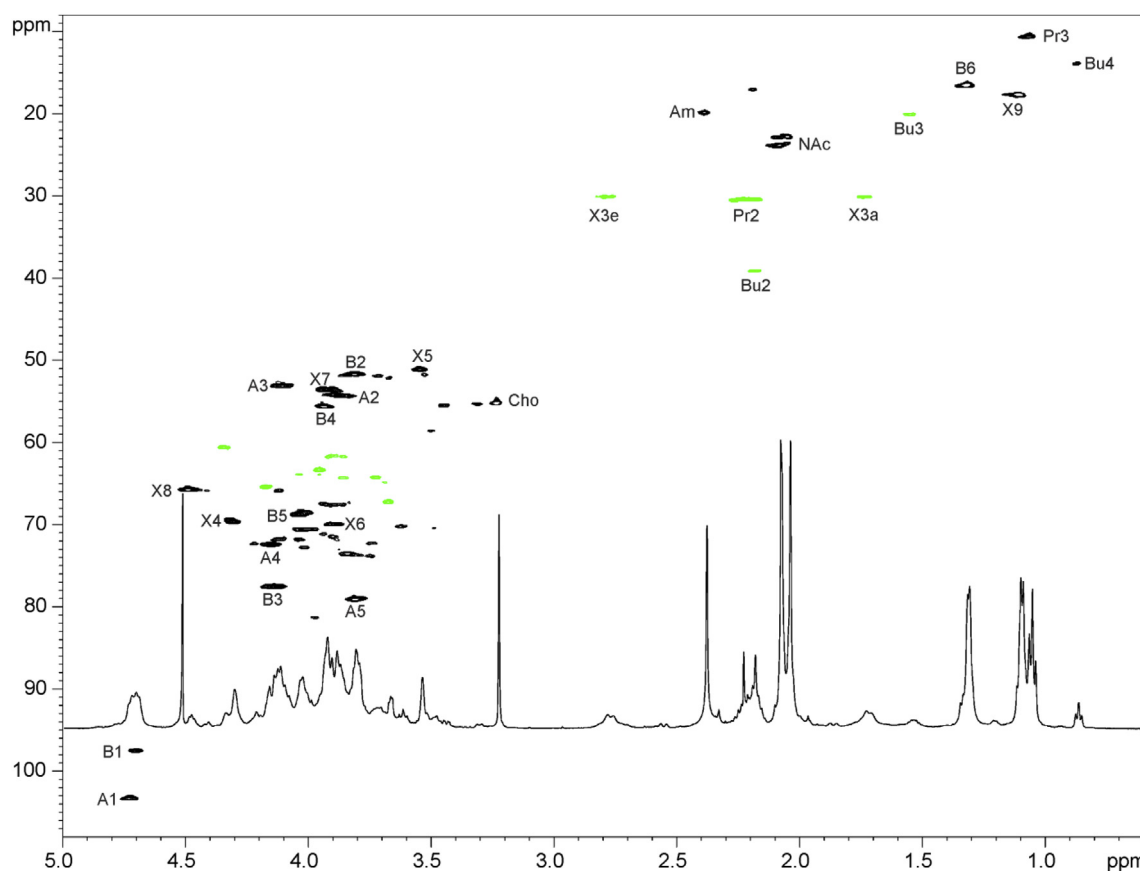
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**Fig. 1.**  $^1\text{H}$ - $^{13}\text{C}$  HSQC and  $^1\text{H}$  NMR spectra of the OPS from *F. nucleatum* ATCC 23726 (600 MHz, 45 °C). Minor signals are from the core oligosaccharide. A, B and X indicate GlcN3NA, FucN4N and Non, respectively; Am, Pr, Bu and Cho indicate acetimidoyl, propanoyl, butanoyl and choline, respectively.

**Table 1**

$^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts ( $\delta$ , ppm) for the OPS, core-RU and OS1 from *F. nucleatum* ATCC 23726. B\* and B\*\* indicate isomeric alditols in the OS1.

Unit		1	2	3 (ax, eq)	4	5	6	7	8	9
X Non core-RU	$^1\text{H}$			1.86, 2.56	4.12	3.52	4.14	3.93	4.54	1.12
	$J_{n,n+1}$		$J_{3eq,4}$ 4.5	$J_{3ax,4}$ 4.5	< 3	< 3	11	< 3	6.5	
	$^{13}\text{C}$	175.6	99.6	36.0	66.0	51.8	69.5	53.7	65.9	17.9
B FucN4N core-RU	$^1\text{H}$	4.51	3.85	4.13	3.94	4.01	1.35			
	$^{13}\text{C}$	103.1	51.9	77.7	55.6	68.6	16.7			
A GlcN3NA core-RU	$^1\text{H}$	4.74	3.88	4.11	4.21	3.81				
	$^{13}\text{C}$	103.5	54.4	53.2	72.5	79.3	176.5			
X Non PS	$^1\text{H}$			1.73, 2.78	4.31	3.54	3.90	3.92	4.48	1.10
	$^{13}\text{C}$			30.1	69.6	51.1	70.0	53.6	65.8	17.7
B FucN4N PS	$^1\text{H}$	4.70	3.81	4.14	3.93	4.03	1.32			
	$^{13}\text{C}$	97.5	51.7	77.6	55.6	68.7	16.5			
A GlcN3NA PS	$^1\text{H}$	4.73	3.86	4.10	4.15	3.81				
	$^{13}\text{C}$	103.3	54.3	53.2	72.4	79.1				
A GlcN3NA OS1	$^1\text{H}$	1.74	3.91	4.20	4.27	4.09				
	$^{13}\text{C}$	102.6	54.4	53.1	72.1	77.4	174.9			
X* Non OS1	$^1\text{H}$			1.84, 2.54	4.00	3.82	4.30	3.85	4.38	1.14
	$^{13}\text{C}$	173.0	98.4	35.5	66.9	48.2	70.4	54.0	66.1	19.1
B* OS1	$^1\text{H}$	3.51; 3.57	3.97	4.02	1.54; 1.73		1.16			
	$^{13}\text{C}$	61.2	55.6	78.0	41.8	65.0	23.6			
B** OS1	$^1\text{H}$	3.51; 3.57	4.05	4.03	1.58; 1.88		1.14			
	$^{13}\text{C}$	61.2	54.2	77.5	41.0	65.3	23.3			
Pr OS1	$^1\text{H}$		2.26; 2.26	1.08						
	$^{13}\text{C}$	179.5	30.2	10.6						

Additional signals in the NMR spectra of the OPS: NAc at 175.2, 2.04/22.8; 175.2, 2.05/23.6 ppm; NAm at 167.2, 2.38/19.8 ppm; NFo at 8.07/171.5 ppm; N-propanoyl at 179.0, 2.05–2.10/30.4, 0.93/10.7 ppm; N-butanoyl at 179.0, 2.05/39.1, 1.43/20.3, 0.74/13.6 ppm.

It had small vicinal coupling constants  $J_{3a,4}$ ,  $J_{3e,4}$ ,  $J_{4,5}$ ,  $J_{5,6}$ ,  $J_{7,8}$  (2–5 Hz), and one large coupling constant  $J_{6,7} = 11$  Hz (Table 1). These data and  $^{13}\text{C}$  NMR chemical shifts (Table 1) agreed with published data for the C-4 – C-7 fragment of the corresponding L-glycero-L-gluco isomer

[11]. However, large deviations were observed for C-8 (~2.5 ppm) and C-9 (~1 ppm) chemical shifts. No data for the D-glycero-L-gluco isomer are available but the C-8 and C-9 chemical shifts of Non fit better to the corresponding fragment in the D-glycero-L-manno isomer as compared

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