

## PRIMARY STRUCTURE OF THE *Klebsiella* SEROTYPE-51 CAPSULAR POLYSACCHARIDE

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### ABSTRACT

The *Klebsiella* serotype-51 (K51) capsular polysaccharide consists of a tetra-saccharide repeating-unit comprising a  $\rightarrow 3$ )- $\alpha$ -D-Galp-(1 $\rightarrow$ 3)- $\alpha$ -D-Galp-(1 $\rightarrow$  chain substituted by an  $\alpha$ -D-GlcpA-(1 $\rightarrow$ 6)- $\alpha$ -D-Glcp-(1 $\rightarrow$  branch at position 4 of one galactosyl residue.

### INTRODUCTION

Of the approximately eighty *Klebsiella* capsular polysaccharides of different K serotypes<sup>1–3</sup>, more than fifty have been subjected to analyses of primary structure (e.g. refs. 3–7). We now report on the structure of the serotype-51 glycan.

### MATERIAL AND METHODS

*Bacteria.* — *Klebsiella* 4715/50 (O3:K51), the serological test-strain for the *Klebsiella* K51 antigen, was kindly supplied by Dr. Ida Ørskov (WHO International Escherichia Center, Statens Seruminstitut, Copenhagen).

All materials and methods used have been described or cited previously<sup>8–10</sup>.

### RESULTS

*Klebsiella* serotype-51 capsular glycan was isolated from *Klebsiella* 4715/50 (O3:K51). The sugar composition and some properties of the material are shown in Table I. The oligosaccharides obtained from the glycan by partial hydrolysis with

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TABLE I

COMPOSITION AND SOME PROPERTIES OF *Klebsiella* SEROTYPE-51 CAPSULAR POLYSACCHARIDE

Polysaccharide	Weight percent of sugar residues (molar ratios) <sup>a</sup>				$s_{20}^c$ (10 <sup>-13</sup> sec)	Equivalent weight <sup>d</sup>	[ $\alpha$ ] <sub>589</sub> <sup>25e</sup> (degrees)
	D-Glc	D-Gal	D-GlcA	Total <sup>b</sup>			
Native <sup>f</sup>	17.2 <sup>g</sup> (1.00)	38.1 <sup>h</sup> (2.22)	19.8 (1.06)	94.5 <sup>b</sup>	1) 3.6 <sup>c</sup> 2) 6.1	645	+95
Alkali-treated <sup>i</sup>	18.5 (1.00)	39.2 (2.12)	20.2 (1.01)	n.d.	3.1 <sup>c</sup>	n.d.	+107
Carboxyl-reduced <sup>j</sup>	31.3 <sup>k</sup> (2.00)	38.6 (2.47)	6.5 (0.38)	n.d.	n.d.	n.d.	n.d.

<sup>a</sup>Hexoses determined by g.l.c. of the alditol acetates<sup>11</sup>, and hexuronic acid by the carbazole-sulfuric acid method<sup>12</sup>. <sup>b</sup>Including 16.3% of water (loss of weight after 24 h at 50° in *vacuo* over P<sub>2</sub>O<sub>5</sub>), 0.3% of protein (determined after hydrolysis with an amino acid analyser), 0.2% of nucleic acid (estimated from the absorption at 260 nm, using yeast RNA as a standard), and 2.6% of sodium (calculated on the basis of the glucuronic acid value). Paper electrophoresis or g.l.c., after hydrolysis<sup>13</sup>, and <sup>1</sup>H-n.m.r. spectroscopy indicated the absence of pyruvate acetal and *O*-acetyl substituents. <sup>c</sup>0.4%. Solutions (w/v) in phosphate-buffered, physiological saline were centrifuged; like other capsular polysaccharides from Enterobacteriaceae<sup>8,14</sup>, the material did not sediment uniformly before mild treatment with alkali (see footnote *i*). <sup>d</sup>Obtained by titration of the acidic form of the glycan<sup>9</sup>. <sup>e</sup>In water (*c* 0.3). <sup>f</sup>As extracted by the phenol-water-cetyltrimethylammonium bromide procedure<sup>8,14</sup>; 100 14-cm nutrient-agar plates yielded 21.5 g of dry bacteria and, from these, 2.2 g (10.2%) of K51 polysaccharide. <sup>g</sup>After optimum hydrolysis (32 h, 100°, 0.5M H<sub>2</sub>SO<sub>4</sub>); about the same value was obtained with fungal D-glucose oxidase<sup>8</sup>. <sup>h</sup>After an optimum hydrolysis time of 6 h; about the same value was obtained with D-galactose dehydrogenase from *Pseudomonas fluorescens*<sup>8</sup>. <sup>i</sup>For 4 h at 56° in 0.25M NaOH<sup>8,14</sup>. <sup>j</sup>Carboxyl-reduced by the method of Taylor *et al.*<sup>15</sup>. <sup>k</sup>About the same value was obtained with fungal D-glucose oxidase<sup>8</sup>.

acid are listed in Table II, and the results of methylation-g.l.c.-m.s. are given in Table III. Table IV summarises the <sup>1</sup>H-n.m.r. data and the structural deductions.

## DISCUSSION

As evidenced by the sugar composition (Table I) and the <sup>1</sup>H-n.m.r. data (Table IV), the *Klebsiella* serotype-51 capsular glycan consists of tetrasaccharide repeating-units, comprising one D-glucuronic acid, one D-glucose, and two D-galactose residues; it is not substituted by pyruvate acetal or by *O*-acetyl groups (Table I).

The data presented in Tables II and IV demonstrate that the component sugars occur in the sequence GlcA→Glc→Gal→Gal or GlcA→Glc→(Gal→)Gal, and that all anomeric configurations are  $\alpha$ .

The results of methylation-g.l.c.-m.s. (Table III) show that GlcA is branch-terminal in the repeating unit of the polysaccharide (column 5), that it substitutes Glc at position 6 (column 1), and that one of the Gal residues is 3- and the other 3,4-di-substituted (branching point in the chain) (columns 4 and 5). From the

TABLE II

ACIDIC OLIGOSACCHARIDES OBTAINED FROM *Klebsiella* SEROTYPE-51 CAPSULAR POLYSACCHARIDE BY PARTIAL HYDROLYSIS WITH ACID

Determination	A2 <sup>a</sup>	A3	A4 <sup>b</sup>
Yield (%)	27	11	n.d.
Molar ratio of sugar components <sup>c</sup>			
D-Glucose	0.9	0.9	0.8
D-Galactose	—	1.0	1.8
D-Glucuronic acid	1.0	1.0	1.0
Reducing-end sugar <sup>d</sup>	Glc	Gal	Gal
Mobility in paper electrophoresis <sup>e</sup> ( <i>M</i> <sub>GlcA</sub> )	0.67	0.54	0.45
<i>R</i> <sub>Glc</sub> in paper chromatography <sup>f</sup>	0.52	0.21	0.08
Enzymic hydrolysis by exo-glycosidases after reduction <sup>15</sup> with carbodi-imide/NaBH <sub>4</sub>			
α-D-Glucosidase <sup>g</sup>	+	+	+
β-D-Glucosidase <sup>h</sup>	—	—	—
α-D-Galactosidase <sup>i</sup> after α-D-glucosidase <sup>j</sup>	n.d.	n.d.	+
β-D-Galactosidase <sup>k</sup> after α-D-glucosidase <sup>j</sup>	n.d.	n.d.	—

<sup>a</sup>A2, aldobiouronic acid; A3, aldotriouronic acid; *etc.* Obtained from the polysaccharide by preparative, paper electrophoresis (see footnote *e*) after 60 min at 100° in 0.5M H<sub>2</sub>SO<sub>4</sub> (or after 45 min at 98° in 2M trifluoroacetic acid). <sup>b</sup>A4 consists of a mixture of the two possible tetrasaccharides in the ratio ~1:3 (see Table III, column 3; Table IV; and Discussion). <sup>c</sup>Hexoses determined by g.l.c. of the alditol acetates<sup>11</sup>, and hexuronic acid by the carbazole-sulfuric acid method<sup>12</sup>. <sup>d</sup>Identified by g.l.c. as the alditol acetate, after reduction with NaBH<sub>4</sub>, hydrolysis, and preparation of the acetylated aldononitriles from the other sugars<sup>16</sup>. <sup>e</sup>In pyridine-acetic acid-water (10:4:86; pH 5.3). <sup>f</sup>Descending p.c. with ethyl acetate-acetic acid-formic acid-water (18:3:1:4). <sup>g</sup>α-D-Glucosidase from yeast, using maltose and cellobiose as controls, and estimating the D-glucose liberated with fungal D-glucose oxidase<sup>8</sup>. <sup>h</sup>β-D-Glucosidase from sweet almonds, with the same controls and detection. <sup>i</sup>α-D-Galactosidase from green coffee-beans, using melibiose and lactose as controls, and determining the D-galactose liberated with D-galactose dehydrogenase from *Ps. fluorescens*<sup>8</sup>. <sup>j</sup>The tetrasaccharides were incubated for 24 h with a large excess of α-D-glucosidase, and the resulting disaccharide fraction was isolated by chromatography on Biogel P2 and then treated with the galactosidase. <sup>k</sup>β-D-Galactosidase from *Escherichia coli*<sup>8</sup> with the same controls and detection as in footnote *i*.

methylation analysis of the aldotriouronic acid (column 2) [*i.e.*, from its structure GlcpA-(1→6)-GlcP-(1→4)-Gal], it can be deduced that the branch in the repeating unit consists of α-D-GlcpA-(1→6)-α-D-GlcP-(1→, and that the chain comprises two →3)-α-D-Galp-(1→ residues, one of which is substituted by the branch at position 4 (see Table IV for a summary). These structural conclusions are confirmed by the methylation results obtained with the mixture of tetraouronic acids, also isolated on partial hydrolysis of the glycan with acid (Table II; Table III, column 3; Table IV, footnote *b*).

TABLE III

IDENTIFICATION<sup>a</sup> AND RATIOS OF *O*-ACETYL-*O*-METHYLAIDITOLS OBTAINED FROM *Klebsiella* SEROTYPE-51 CAPSULAR POLYSACCHARIDE AND ITS DERIVATIVES

<i>Per-O-acetyl derivative of<sup>b</sup></i>	1 <sup>c</sup>	2	3	4	5
	<i>Ratio of peak integrals<sup>d</sup></i>				
2,3,4,6-GlcOH <sup>e</sup>	—	—	—	—	0.8 <sup>e</sup>
2,3,4,6-GalOH	—	—	1.0	—	—
2,4,6-GalOH	—	—	0.4	1.0	1.0
2,3,6-GalOH	—	1.2	0.4	—	—
2,3,4-GlcOH	1.0	1.0	1.1	1.0	1.1
2,6-GalOH	—	—	1.3	1.1	1.1

<sup>a</sup>By combined gas chromatography-mass spectrometry employing columns packed with ECNSS-M17,<sup>18</sup> and capillary columns wall-coated with Silar 9CP or OV-101<sup>10</sup>. <sup>b</sup>2,3,4,6-GlcOH = 2,3,4,6-tetra-*O*-methyl-D-glucitol, *etc.* <sup>c</sup>1, aldobiouronic acid A2 (see Table II), permethylated; 2, aldotriouronic acid A3, permethylated; 3, mixture of the two possible tetraouronic acids (ratio ~1:3; see Table IV, footnote *b*, and Discussion), permethylated; 4, K51 polysaccharide, permethylated; 5, K51 polysaccharide, permethylated, carboxyl-reduced/dideuterated with calcium borodeuteride<sup>19</sup>, and remethylated. <sup>d</sup>Detection by flame ionisation. <sup>e</sup>Dideuterated at position 6.

TABLE IV

<sup>1</sup>H-N.M.R. SIGNALS OF ANOMERIC PROTONS IN *Klebsiella* SEROTYPE-51 OLIGOSACCHARIDES AND CAPSULAR POLYSACCHARIDE<sup>a</sup>, AND SUMMARY OF STRUCTURES

<i>Oligo- or poly-saccharide</i>	$\delta$	J (Hz)	<i>Approximate ratio of integrals</i>	<i>Proton assignment</i>
Aldobiouronic acid A2	5.20	~3	0.5	6)- $\alpha$ -Glc
$\alpha$ -D-GlcpA-(1 $\rightarrow$ 6)-D-Glc	4.90	~3	1.0	$\alpha$ -GlcpA
	4.63	7.5	0.5	6)- $\beta$ -Glc
Aldotriouronic acid A3	5.29	3.0	0.5	4)- $\alpha$ -Gal
$\alpha$ -D-GlcpA-(1 $\rightarrow$ 6)- $\alpha$ -D-Glcp-(1 $\rightarrow$ 4)-D-Gal	4.96	~2.5	2.0	{ 6)- $\alpha$ -Glcp $\alpha$ -GlcpA
	4.96	~2.5		
	4.63	7.0		
Polysaccharide	5.35	~3	1.0	{ 3)- $\alpha$ -Galp 3,4)- $\alpha$ -Galp
$\alpha$ -D-GlcpA	5.20	~3	1.0	
1	5.08	~3	1.0	
↓	4.98	~3	1.0	
6				
$\alpha$ -D-Glcp				
1				
↓				
4				
$\rightarrow$ 3)- $\alpha$ -D-Galp-(1 $\rightarrow$ 3)- $\alpha$ -D-Galp-(1 $\rightarrow$				
$\uparrow^b$ (1) <sup>b</sup>				

<sup>a</sup>Solutions (0.5–3.0%, w/v) in deuterium oxide were run at 70° and 90 MHz<sup>9</sup>. <sup>b</sup>Formation of tetraouronic acids during partial, acid hydrolysis; preferential and minority cleavage-site (see Table III, column 3, and Discussion).

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