



# Synthesis of the Repeating Unit of the O-specific Polysaccharide of *Shigella sonnei* and Quantitation of its Serologic Activity

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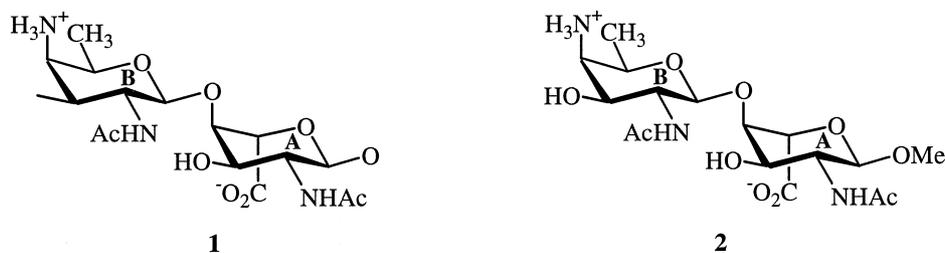
**Abstract**—The chemical synthesis of the zwitterionic disaccharide **2** is described that corresponds to the repeating unit of the O-specific polysaccharide (**1**) of the Gram-negative human pathogen *Shigella sonnei*. Passive hemolysis inhibition tests using a hyper-immune rabbit serum raised against *S. sonnei* showed that the serologic activity of the disaccharide **2** is nearly 2- to 3-fold higher than those of its component monosaccharides. NMR data of **2** are in support of the proposed structure of the O-specific polysaccharide. © 1999 Elsevier Science Ltd. All rights reserved.

*Shigella sonnei* is a Gram-negative bacterium that can cause diarrhea and dysentery in humans. A major cell surface component of this bacterium is its O-specific polysaccharide<sup>1,2</sup> (O-SP) that is the outermost, sero-determinant domain of the highly complex lipopolysaccharide. The O-SP is an essential virulence factor of Shigellae including *S. sonnei*: only strains that have their O-SP fully expressed are virulent. Additionally, the O-SP is related to host immunity in that protection against infection is a correlate of the IgG antibody level against the O-SP.<sup>3</sup> The O-SP of *S. sonnei* is a non-immunogenic molecule (MW ~25 kDa) that must be conjugated to an immunogenic protein carrier to induce antibodies. Clinical trials have confirmed that protein conjugates of the O-SP elicited IgG levels in adult volunteers that were similar to those in patients convalescent from shigellosis.<sup>4</sup> The O-SP of *S. sonnei* is composed of the zwitterionic disaccharide repeating unit **1** consisting of the rare sugars 2,4,6-trideoxy-2-acetamino-4-amino-D-galactose and 2-acetamido-2-deoxy-L-altruronic acid that are connected through 1,2-*trans* interglycosidic linkages.<sup>1,2</sup>

Our approach to vaccine development is based on the assumption that fragments of O-SPs mimic epitopes of the native polymers and thus may induce polysaccharide-specific antibodies when conjugated to an immunogenic carrier. Recent data confirmed this assumption and showed that albumin conjugates of chemically synthesized extended oligosaccharide fragments<sup>5,6</sup> of the O-SP of *S. dysenteriae* type 1 elicited higher levels of the homologous O-SP-specific IgG antibodies than did the conjugate of the native polysaccharide.<sup>7</sup>

As a prelude to chemical synthesis of oligosaccharides corresponding to **1** we have reported the synthesis of the methyl glycosides of its component monosaccharides in their natural glycosidic configuration.<sup>8</sup> We have also presented evidence that in the native polysaccharide both pyranose residues are preferentially in the <sup>4</sup>C<sub>1</sub> chair conformation and that the polysaccharide exists in the zwitterionic form.<sup>9</sup> Here we describe the first synthesis of a disaccharide part (**2**) of the O-SP that to our knowledge is also the first report of a zwitterionic disaccharide containing free amino and carboxyl groups at alternating monosaccharide residues. The disaccharide **2** was constructed from an appropriately protected altruronic acid derivative (**5**) that was used as the acceptor and from the activated donor **9**. The common

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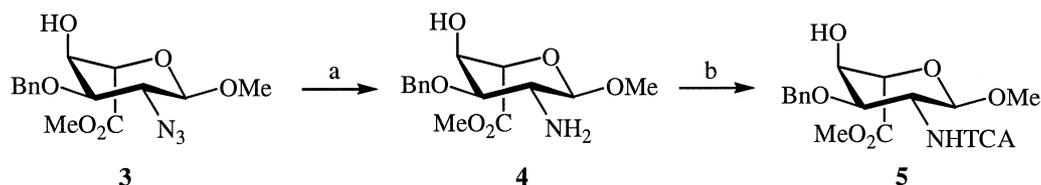
feature of both intermediates is the presence of a trichloroacetyl protecting group at their respective amino function. This protecting group has previously been shown to be suitable for *N*-protection without adverse outcomes during glycosylation reactions.<sup>10</sup> It was also demonstrated to have good *trans*-directing effect and to provide the substrates with acceptable solubility properties. The synthetic steps to the key donor and acceptor intermediates are described next.

**The altruronic acid residue 5 (Scheme 1).** The synthetic route started from the glycoside **3** that was available from an earlier work.<sup>8</sup> In the first step the azido group in compound **3** was reduced to an amino group with H<sub>2</sub> over Pd/C to afford **4**. Hydrogenolytic cleavage of the *O*-benzyl group in this step was prevented by the use of one equivalent of pyridine. Subsequent treatment of **4** with trichloroacetyl chloride afforded **5** in 56% overall yield.

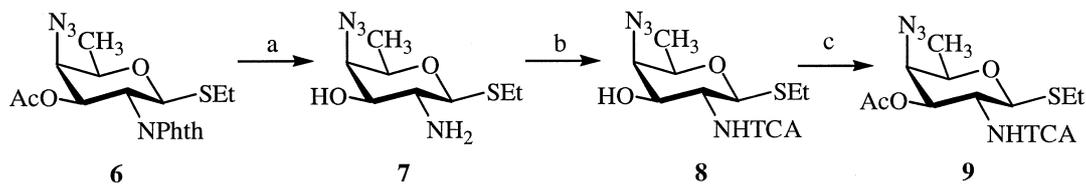
**The trideoxy-galactose residue 9 (Scheme 2).** Starting compound was thioglycoside **6** that was converted to the amine **7** by treatment with ethylenediamine, in 94%

yield. Next, a trichloroacetyl group was installed at the free amino group by exposure of **7** to trichloroacetyl chloride in the presence of triethylamine ( $\rightarrow$ **8**). The synthesis of the glycosyl donor **9** was completed by acetylation of **8** with pyridine/acetic anhydride, in 94% overall yield for two steps.

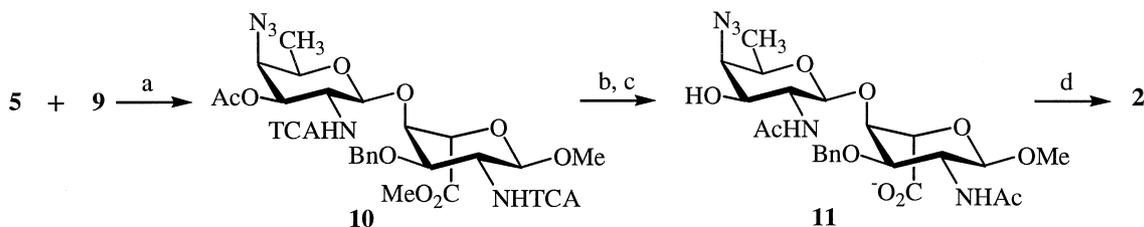
Having readied the donor (**9**) and the acceptor moieties (**5**), the fully protected disaccharide **10** was prepared by NIS/TfOH-promoted activation<sup>11,12</sup> of the thioglycoside **9** in the presence of 0.55 equiv amount of the acceptor **5** in 85% yield (Scheme 3). Replacement of the *N*-trichloroacetyl groups with the acetyl groups was previously described through Bu<sub>3</sub>SnH-mediated reductive cleavage of the chlorine atoms.<sup>10</sup> This approach failed to give satisfactory results when applied to **10**. Fortunately, treatment of **10** with NaOH in MeOH excellently removed the *N*-trichloroacetyl groups to afford an intermediate that was converted to the *N*-acetamido derivative **11** by conventional acetylation, in 72% overall yield for two steps. As the final phase in the synthetic sequence, compound **11** was subjected to



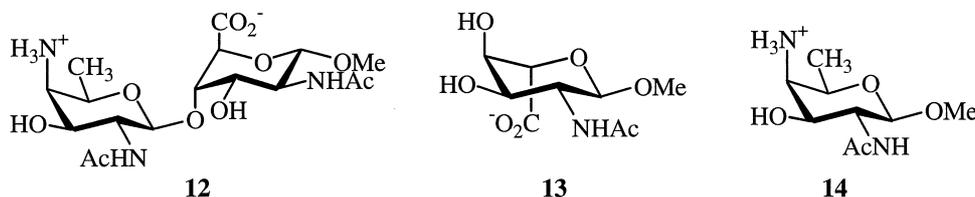
**Scheme 1.** Reagents and conditions: (a) H<sub>2</sub>/Pd-C, MeOH, C<sub>5</sub>H<sub>5</sub>N, 25 °C, 2 h 81%; (b) 1.1 equiv of Cl<sub>3</sub>CCOCl, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 0 °C, 1 h 70%.



**Scheme 2.** Reagents and conditions: (a) NH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>NH<sub>2</sub> (excess), EtOH, reflux, 2 h, 94%; (b) 1.2 equiv of Cl<sub>3</sub>CCOCl, Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, 0 °C, 20 min; (c) C<sub>5</sub>H<sub>5</sub>N, Ac<sub>2</sub>O, 25 °C, 2 h, 94% for two steps.



**Scheme 3.** Reagents and conditions; (a) 1.8 equiv of **9**, 2.3 equiv of NIS, TfOH (cat), CH<sub>2</sub>Cl<sub>2</sub>, -30 °C, 1.5 h, 85%; (b) aq NaOH, MeOH, 25 °C, 2.5 days; (c) Ac<sub>2</sub>O, MeOH, 0 °C, 50 min, 72% for two steps; (d) H<sub>2</sub>/Pd(OH)<sub>2</sub>-C, EtOH, AcOH 25 °C, 4 days, 51%.



**Table 1.** Inhibition of hemolysis of *Shigella sonnei* lipopolysaccharide-sensitized erythrocytes by the lipopolysaccharide-related antigens **2**, **13** and **14**<sup>a</sup>

Antigen	IC <sub>50</sub> (mM) <sup>b</sup>
<b>2</b>	3.9
<b>13</b>	6.9
<b>14</b>	9.8

<sup>a</sup>IC<sub>50</sub>: 50% inhibition concentration.

<sup>b</sup>Estimated range of error: ≤10%.

hydrogenolytic cleavage of the *O*-benzyl groups that simultaneously reduced the azido group to afford the targeted disaccharide **2**.<sup>13</sup>

A comparison of the <sup>1</sup>H and <sup>13</sup>C NMR chemical shifts for compounds **1**,<sup>9</sup> and **2** shows an excellent agreement of the corresponding resonances that lends further support for the absolute configuration of the altruronic residue as being L and for the trideoxy-galactose residue as being D.<sup>14</sup> For a comparison, we have also synthesized the diastereomeric disaccharide **12** that contains the enantiomeric, D altruronic acid moiety.<sup>15</sup> For compound **12** the chemical shift of the C-5 carbon atom of the uronic acid residue is 70.5 ppm, distinctly different from the corresponding resonances in **1** and **2** (78.4 and 76.9, respectively). Minor differences in the chemical shifts for residues A in **1** and **2** are likely due to conformational differences and indicate that the disaccharide can only partially express conformational features of polysaccharide **1**.

The antigenicity of the disaccharide **2** was assayed by the passive hemolysis inhibition test as described by Kontrohr and Péterffy.<sup>16</sup> The method is based on complement-mediated hemolysis of lipopolysaccharide-sensitized erythrocytes after anti-LPS antibody binding. Using sheep erythrocytes, guinea pig complement, phase I lipopolysaccharide of *S. sonnei*,<sup>1</sup> and hyper-immune polyclonal rabbit serum raised against *S. sonnei*, the concentration of the disaccharide **2** and those of the component monosaccharides **13** and **14** were determined that were necessary for 50% inhibition of hemolysis. The results (Table 1) show that the disaccharide **2** is a better inhibitor than either one of its monosaccharide components of which the altruronic acid derivative **13** is superior. Less than 20% inhibition could be observed with the unnatural disaccharide **12** up to 15 mM concentration and no inhibition was seen with unrelated saccharides.

In summary, we have synthesized for the first time the complete repeating unit of the O-specific polysaccharide

of *Shigella sonnei* and presented spectroscopic and immunochemical support for the structure of the native polysaccharide. Current work is directed at the synthesis and immunochemical study of the frame-shifted, alternative disaccharide repeating unit and higher-membered saccharides corresponding to the O-specific polysaccharide of *S. sonnei* for their eventual use as vaccine-components against enteric diseases.

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- All new compounds had satisfactory analytical, <sup>1</sup>H, <sup>13</sup>C NMR, and mass spectral data. For **2** <sup>1</sup>H NMR(D<sub>2</sub>O): δ 4.73 (H-1<sub>B</sub>), 4.70 (H-1<sub>A</sub>), 4.55 (H-5<sub>A</sub>), 4.42 (H-4<sub>A</sub>), 3.90 (H-3<sub>B</sub>), 3.88 (H-2<sub>A</sub>), 3.88 (H-5<sub>B</sub>), 3.81 (H-2<sub>B</sub>), 3.70 (H-3<sub>A</sub>), 1.31 (H-6<sub>B</sub>); <sup>13</sup>C NMR: δ 102.9 (C-1<sub>B</sub>), 100.3 (C-1<sub>A</sub>), 77.9 (C-4<sub>A</sub>), 76.9 (C-5<sub>A</sub>), 69.3 (C-3<sub>B</sub>), 69.2 (C-3<sub>A</sub>, C-5<sub>B</sub>), 54.6 (C-4<sub>B</sub>), 52.8 (C-2<sub>B</sub>), 52.1 (C-2<sub>A</sub>), 16.3 (C-6<sub>B</sub>).
- We had no specific reason to question the reported structure of the O-SP of *S. sonnei*. However, in the light of numerous structural revisions of bacterial polysaccharides it is always advisable to obtain independent structural proof for these materials before undertaking a synthetic project towards their extended portions.
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