



## **Brucellosis Detection**



## Design and Synthesis of a Universal Antigen to Detect Brucellosis\*\*

Julie Guiard, Eugenia Paszkiewicz, Joanna Sadowska, and David R. Bundle\*

In memory of Malcolm Perry

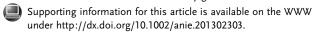
Brucellosis is a highly contagious zoonosis primarily caused by ingestion of unsterilized milk or meat from animals infected by members of the genus Brucella.[1] Bovine brucellosis is caused by B. abortus, less frequently by B. melitensis, and occasionally by B. suis. B. abortus in cattle causes premature abortion in pregnant cows. In humans, Brucella species cause a nonfatal but debilitating disease. Infection is widespread globally and in those parts of the world where malaria is endemic it would be useful to have a simple diagnostic test to differentiate between malaria and Brucella or other microbes as the cause of febrile fever. [2,3] Presumptive diagnosis depends on detection of antibodies to Brucella A and M antigens and is confirmed by microbiological culture. [4] The A- and M-antigenic determinants are expressed simultaneously on the O-antigen polysaccharide domain of Brucella smooth lipopolysaccharides (sLPS), and this sLPS is used to detect antibodies present in sera of animals or humans suspected of being infected. Brucella is a virulent pathogen requiring level 3 containment, rendering the production of diagnostic O-antigens a demanding and specialized task. An alternative diagnostic, the O-antigen of Yersinia enterocolitica O:9 (level 2 containment) is an exclusively A antigen devoid of M epitopes.

Brucella sLPS is resistant to partial degradation methods that would permit isolation of pure A- or M-antigenic determinants in quantities for practical application. A well-defined synthetic antigen that incorporates both A and M epitopes combined with a versatile tether that allows attachment to surfaces, particles, and polymers could replace the native O-antigen in diagnostic tests, including those required for simple, on-site tests in remote locations.<sup>[5]</sup>

The O-antigen of *Brucella* sLPS is a homopolymer of the rare sugar 4,6-dideoxy-4-formamido-α-D-mannose (α-D-Rha4NFo). Three *Brucella* antigenic phenotypes A<sup>+</sup>M<sup>-</sup>, A<sup>-</sup>M<sup>+</sup>, and A<sup>+</sup>M<sup>+</sup> were characterized using NMR spectroscopy and monoclonal antibodies and confirmed that a single O-polysaccharide molecule incorporates both A-and M-antigenic determinants (Figure 1). The A epitope is dominant in most strains of *B. abortus* while the M epitope is characteristic of *B. melitensis*. Prototypical A structures



<sup>[\*\*]</sup> This work was made possible by a Natural Science and Engineering Research Council of Canada Discovery grant to D.R.B.



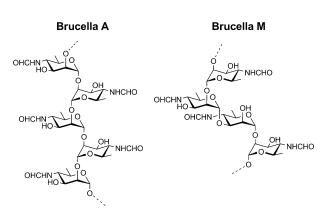


Figure 1. Structures of the A and M antigenic determinants of B. abortus and B. melitensis, respectively.

consist predominantly of  $\alpha$ 1,2-linked D-Rha4NFo residues. [6a,7b] In its highly expressed form, the M-antigenic determinant was first proposed to contain one  $\alpha$ 1,3 linkage for every four  $\alpha$ 1,2-linked residues. [6c,7b] A revised structure was reported, where the ratio of 1,2 to 1,3 linkages is 3:1. [7c]

Even in *Brucella* strains defined as  $A^+M^-$ , an  $\alpha 1,3$ -linked D-Rha4NFo is present and generally occurs at least once in every 50 residues. This irregular structural feature is consistent with the biosynthesis of these polysaccharides by the ABC-type translocation pathway. Two genes, wzm and wzt, which typically form the ABC transporter/exporter of the ABC-2 subfamily, are present in the *B. melitensis* genome. [9] The precise structure of the M-antigenic determinant which is devoid of any A activity remains uncertain.

Our work with monoclonal antibodies and synthetic oligomers established that the antigenic determinant of the *B. abortus* A antigen is most likely a tetrasaccharide of contiguous  $\alpha 1,2$ -linked D-Rha4NFo residues. [6e,7b] While the M antigen could in principle be characterized as an  $\alpha 1,3$ -linked D-Rha4NFo disaccharide, this antigenic determinant is more likely to be defined as a larger oligosaccharide with adjacent  $\alpha 1,2$ -linked D-Rha4NFo residues that are sufficiently short to preclude recognition by A-specific antibodies. [7b]

Pentasaccharide 1 (Figure 2) was selected as the largest sized antigen that might selectively exhibit M-type characteristics with limited cross reaction with A-specific antibodies. Nonasaccharide 2 contains two A- and one M-type epitopes (Figure 2) and should serve as a universal antigen to detect antibodies in animals or humans infected by *B. abortus*, *B. melitensis*, and *B. suis*.

The large size of the target oligosaccharides **1** and **2**, and the incorporation of an internal 1,3 linkage and a tether for



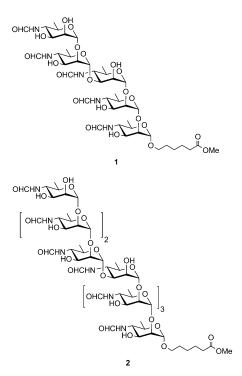


Figure 2. Target pentasaccharide 1 anticipated to exhibit preferred binding to M-specific antibodies and nonasaccharide 2 designed to bind both A- and M-specific antibodies.

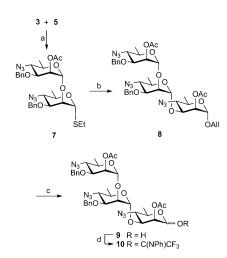
antigen synthesis had not been previously attempted<sup>[10]</sup> and necessitated the development of an improved synthetic strategy. The linker 5-methoxycarbonylpentanol<sup>[11,12]</sup> is compatible with the deprotection of the assembled oligosaccharides and opens up several routes for subsequent conjugation to protein to create glycoconjugate antigens. [11,13]

Monosaccharide synthons 3-6 required for the assembly of 1 and 2 were accessed by published methods (Scheme 1).<sup>[10]</sup> N-phenyl trifluoroacetimidates were employed for glycosylation reactions since this donor has been shown to be more efficient than the corresponding trichloroacetimidate deriva-

Scheme 1. Key synthons for assembly of penta- and nonasaccharides 1 and 2. All = all vl.

tive for glycosylations involving 6-deoxy sugar donors.<sup>[14]</sup> Glycosylation of 5 by 3 was performed in the presence of trimethylsilyl trifluoromethanesulfonate (TMSOTf) to give disaccharide 7 in 94% yield with complete stereocontrol and without detectable amounts of the  $\beta$ -anomer (Scheme 2).

The 1,3-linked trisaccharide building block 8 was created as its allyl glycoside to allow facile access to a hemiacetal and subsequently an imidate leaving group. [15] Glycosylation reactions were tried with allyl as a leaving group but all attempts failed. [16] Consequently, 8 was selectively depro-



**Scheme 2.** a) TMSOTf,  $CH_2Cl_2$ , 94%; b) 6, NIS, TfOH,  $CH_2Cl_2$ , -30°C, 65%; c) PdCl<sub>2</sub>, AcONa, AcOH, H<sub>2</sub>O, 73%; d) CF<sub>3</sub>C(NPh)Cl, Cs<sub>2</sub>CO<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 79%. NIS = N-iodosuccinimide, TfOH = trifluoromethanesulfonic acid.

tected with palladium chloride in acetic acid[15] to give hemiacetal 9, which was converted to the N-phenyl trifluoroacetimidate donor 10 (Scheme 2).

Thioglycoside 7 gave direct access to hemiacetal 11 and subsequently imidate 12, which was used to glycosylate the six carbon linker 13.[12b] Better yields and selectivity in the preparation of 14 were obtained by performing the reaction in toluene at 100°C.[17] Transesterification of 14 gave the tether glycoside acceptor 15 (Scheme 3).

With the two building blocks, trisaccharide 10 and disaccharide glycoside 15, in hand, we obtained pentasaccharide 16 in 68% yield using TMSOTf as the activator (Scheme 4). Stepwise deprotection followed the sequence: deacetylation to give 17 in quantitative yield, followed by reduction of the azido group with hydrogen sulfide to give 18. The amine 18 was directly formylated by a mixed anhydride (acetic anhydride/formic acid 2:1) to give 19.[10b] Following

**Scheme 3.** a) NIS, H<sub>2</sub>O, Me<sub>2</sub>CO, 76%; b) CF<sub>3</sub>C(NPh)Cl, Cs<sub>2</sub>CO<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 90%; c) TMSOTf, toluene, 100°C, 53%; d) MeONa, MeOH,

7323



**Scheme 4.** a) TMSOTf,  $CH_2Cl_2$ , 68%; b) MeONa, MeOH, 86%; c)  $H_2S$ ,  $Py/Et_3N$  1:1; d)  $Ac_2O/HCOOH$  2:1, MeOH, 63% from **17**; e)  $H_2$ , Pd/C, AcOH, 54%. Py = pyridine.

introduction of the N-formamido groups, NMR analyses of all subsequent compounds became difficult due to the presence of E/Z rotamers for each formyl group, leading to a potential mixture of 32 isomers. Their identity was confirmed by a limited set of characteristic NMR resonances and high-resolution mass measurements. Pentasaccharide 1 was obtained by hydrogenolysis of benzyl ethers.

Synthesis of nonasaccharide **2** was envisaged as the creation of a pentasaccharide donor terminated by a 1,3 linkage which would then allow for a pentasaccharide donor with a participating group at C-2 to guide the stereoselective  $\alpha$ -glycosylation of an exclusively 1,2-linked tetrasaccharide. To achieve the synthesis of the pentasaccharide donor, compound **7** was deprotected to give the corresponding acceptor **20**, which was glycosylated by imidate donor **12** as described for **14**. Tetrasaccharide thioglycoside **21** was used directly as the donor for glycosylation of the monosaccharide glycoside **6** to give the  $\alpha$ 1,3-linkage. The allyl group of pentasaccharide **22** was then removed to give **23** and the imidate donor **24** was obtained following reaction with *N*-phenyl trifluoroacetimidoyl chloride (Scheme 5).

Tetrasaccharide tether glycoside **25** was obtained by a 2+2 glycosylation of disaccharide acceptor **15** by the disaccharide donor **12**. Transesterification of **25** gave the tetrasaccharide acceptor **26** which was glycosylated by pentasaccharide donor **24** to give nonasaccharide **27** in 30% yield. The low yield of the 5+4 glycosylation reaction was attributed to the low reactivity of the acceptor. Paucity of the oligosaccharide reactants prevented optimization of this reaction step. The sequence of deprotection steps (deacetylation to **28**, reduction of the azide to **29**, *N*-formylation to **30**, and hydrogenation) to give **2** followed the order used to obtain pentasaccharide **1** (Scheme 6).

The final compounds 1 and 2 were purified by reversephase HPLC. Full NMR assignments were performed on the azido penta- and nonasaccharide derivatives 17 and 28. Selected characteristic NMR resonances and high-resolution mass spectra confirmed the identity of derivatives 19 and 30 and the target oligosaccharides 1 and 2.

**Scheme 5.** a) MeONa, MeOH,81%; b) **12**, TMSOTf, toluene, 100°C, 81%; c) **6**, NIS, TfOH, CH<sub>2</sub>Cl<sub>2</sub>, 68%; d) PdCl<sub>2</sub>, AcONa, AcOH, H<sub>2</sub>O, 62%; e) CF<sub>3</sub>C(NPh)Cl, Cs<sub>2</sub>CO<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 79%.

12 + 15

a

$$N_3$$
BnO

 $N_3$ 
BnO

**Scheme 6.** a) TMSOTf, toluene,  $100^{\circ}$ C, 77%; b) MeONa, MeOH, 84%; c) **24**, TMSOTf,  $CH_2CI_2$ , 30%; d) MeONa, MeOH, 78%; e)  $H_2S$ , Py/Et<sub>3</sub>N 1:1; f) Ac<sub>2</sub>O/HCOOH 2:1, MeOH, 62% from **28**; g)  $H_2$ , Pd/C, AcOH, 48%.



The linker ester moieties of the pentasaccharide and nonasaccharide glycosides 1 and 2 were converted to the respective amides by reaction with ethylenediamine. The amides were reacted with dibutyl squarate and the half esters isolated by reverse-phase HPLC (Scheme 2; see the Supporting Information). The pentasaccharide and nonasaccharide bovine serum albumin (BSA) glycoconjugates were prepared by reaction of a twenty to one molar ratio of the two half esters with BSA. MALDI-TOF mass spectrometry indicated that each conjugate contained approximately 16 copies of the oligosaccharides per molecule of BSA.

Two monoclonal antibodies (YsT9-1 and Bm10) specific for the Brucella A and M antigens[7b] were titred to their end point against the pentasaccharide and nonasaccharide antigens coated on ELISA plates (Figure 3). The nonasaccharide antigen binds A- and M-specific antibodies with equivalent avidity, whereas the pentasaccharide displays a preference for

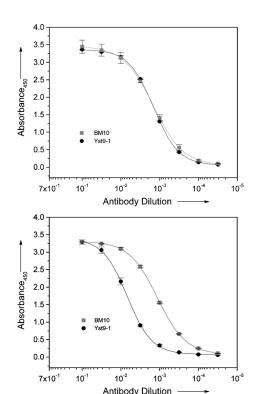


Figure 3. ELISA titration curves for Brucella A- and M-specific mAbs, YsT9-1 and Bm10; pentasaccharide conjugate (top) and nonasaccharide conjugate (bottom).

the M-specific antibody, while still binding the A-specific antibody but with an approximately 10-fold reduced avidity.

Drawing on inferences from studies of monoclonal antibody (mAb) binding specificities, [7b] we have designed unique antigens that are not available from natural sources thereby creating a novel, universal antigen to detect Brucella antibodies that arise during infection by all Brucella strains producing a sLPS. This provides a valuable and convenient antigen for presumptive diagnosis and one that can be deployed in virtually any assay format including those that do not require sophisticated equipment, unavailable in remote locations.[5]

The nonasaccharide glycoconjugate binds Brucella A (YsT9-1) and M (Bm10) mAbs with equal avidity even though these antibodies possess avidity differences of between 400-1000 for the respective A and M O-polysaccharide antigens.<sup>[7b]</sup> The pentasaccharide antigen shows a preference for M-specific mAb. This discrimination between M and A antibodies by synthetic conjugates is expected to improve by decreasing the number of 1,2-linked α-D-Rha4NFo residues from the three present in pentasaccharide 1 to two or one in related tetra- or trisaccharides. By systematically shortening the length of nonasaccharide 2 it should also be possible to arrive at an even shorter oligosaccharide capable of binding both A and M antibodies.

Received: March 18, 2013 Published online: June 11, 2013

**Keywords:** antigens · brucellosis · glycans · polysaccharides

- [1] M.-L. Boschiroli, V. Foulongne, D. O'Callaghan, Curr. Opin. Microbiol. 2001, 4, 58-64.
- [2] J. A. Crump, F. G. Youssef, S. P. Luby, M. O. Wasfy, J. M. Rangel, M. Taalat, S. A. Oun, F. J. Mahoney, Emerging Infect. Dis. 2003, 9,539-544.
- [3] M. D. Ari, A. Guracha, M. A. Fadeel, C. Njuguna, M. K. Njenga, R. Kalani, H. Abdi, O. Warfu, V. Omballa, C. Tetteh, R. F. Breiman, G. Pimentel, D. R. Feikin, Am. J. Trop. Med. Hyg. **2011**, 85, 909-912.
- [4] M. J. Corbel, "Brucellosis in humans and animals". WHO/CDS/ EPR/2006.7.
- [5] a) A. W. Martinez, S. T. Phillips, G. M. Whitesides, Proc. Natl. Acad. Sci. USA 2008, 105, 19606-19611; b) C.-M. Cheng, A. W. Martinez, J. Gong, C. R. Mace, S. T. Phillips, E. Carrilho, K. A. Mirica, G. M. Whitesides, Angew. Chem. 2010, 122, 4881-4884; Angew. Chem. Int. Ed. 2010, 49, 4771-4774.
- [6] a) M. Caroff, D. R. Bundle, M. B. Perry, J. W. Cherwonogrodzky, J. R. Duncan, Infect. Immun. 1984, 46, 384-388; b) D. R. Bundle, J. W. Cherwonogrodzky, M. Caroff, M. B. Perry, Ann. Inst. Pasteur/Microbiol. 1987, 138, 92-98; c) D. R. Bundle, J. W. Cherwonogrodzky, M. B. Perry, Biochemistry 1987, 26, 8717-8726; d) J. W. Cherwonogrodzky, M. B. Perry, D. R. Bundle, Can. J. Microbiol. 1987, 33, 979 – 981; e) P. J. Meikle, M. B. Perry, J. W. Cherwonogrodzky, D. R. Bundle, Infect. Immun. 1989, 57, 2820 - 2828.
- [7] a) D. R. Bundle, M. A. J. Gidney, M. B. Perry, J. R. Duncan, J. W. Cherwonogrodzky, Infect. Immun. 1984, 46, 389-393; b) D. R. Bundle, J. W. Cherwonogrodzky, M. A. Gidney, P. J. Meikle, M. B. Perry, T. Peters, Infect. Immun. 1989, 57, 2829 – 2836; c) J. Kubler-Kielb, E. Vinogradov, Carbohydr. Res. 2013, DOI: 10.1016/j.carres.2013.03.021.
- [8] D. Bronner, B. R. Clarke, C. Whitfield, Mol. Microbiol. 1994, 14, 505 - 519.
- [9] F. Godfroid, A. Cloeckaert, B. Taminiau, I. Danese, A. Tibor, X. de Bolle, P. Mertens, J.-J. Letesson, Res. Microbiol. 2000, 151, 655 - 668.
- [10] a) T. Peters, D. R. Bundle, J. Chem. Soc. Chem. Commun. 1987, 1648-1650; b) D. R. Bundle, M. Gerken, T. Peters, Carbohydr. Res. 1988, 174, 239-251; c) T. Peters, D. R. Bundle, Can. J. Chem. 1989, 67, 491-496; d) T. Peters, D. R. Bundle, Can. J. Chem. 1989, 67, 497-502.

7325



- [11] R. U. Lemieux, D. R. Bundle, D. A. Baker, J. Am. Chem. Soc. 1975, 97, 4076–4083.
- [12] a) G. Solladié, C. Ziani-Cherif, J. Org. Chem. 1993, 58, 2181 –
   2185; b) S. El Fangour, A. Guy, V. Despres, J.-P. Vidal, J.-C. Rossi, T. Durand, J. Org. Chem. 2004, 69, 2498 2503.
- [13] V. P. Kamath, P. Diedrich, O. Hindsgaul, *Glycoconjugate J.* 1996, 13, 315–319.
- [14] a) E. Bedini, A. Carabellese, M. Schiattarella, M. Parrilli, Tetrahedron 2005, 61, 5439-5448; b) D. Comegna, E. Bedini,
- A. Di Nola, A. Iadonisi, M. Parrilli, *Carbohydr. Res.* **2007**, *342*, 1021–1029; c) S. Hanashima, B. Castagner, D. Esposito, T. Nokami, P. H. Seeberger, *Org. Lett.* **2007**, *9*, 1777–1779.
- [15] Y. Du, M. Zhang, F. Kong, Tetrahedron 2001, 57, 1757-1763.
- [16] P. Wang, P. Haldar, Y. Wang, H. Hu, J. Org. Chem. 2007, 72, 5870–5873.
- [17] S.-J. Hou, P. Kováč, Carbohydr. Res. 2010, 345, 999–1007.