O-chain expression in the rough Brucella melitensis strain B115: induction of O-polysaccharide-specific monoclonal antibodies and intracellular localization demonstrated by immunoelectron microscopy

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Spleen cells from mice infected with the rough Brucella melitensis strain B115 were fused with NSO myeloma cells. Hybridoma supernatants were screened in ELISA with cell walls (CW), sonicated cell extracts (CE) and rough lipopolysaccharide (R-LPS) of B. melitensis strain B115 and whole B. melitensis B115 cells. Surprisingly, 22 monoclonal antibodies (mAbs) reacting in ELISA with both CW and CE but not with R-LPS and bacterial cells were shown by immunoblot analysis and ELISA to react with smooth lipopolysaccharide (S-LPS). These mAbs also reacted in ELISA with O polysaccharides (OPS) from the smooth Brucella abortus strain 99 and the smooth B. melitensis strain 16M and thus recognize epitopes present on the O-chain. Proteinase K LPS preparations from B. melitensis B115 analysed by immunoblotting with one mAb (12G12) recognizing S-LPS of both A and M specificity displayed the typical S-LPS high-molecular-mass ladder pattern but no S-LPS was detected in the phenol/water/chloroform/light petroleum LPS preparation of the same strain. mAb 12G12, specific for S-LPS, and a mAb (A68/03F03/D05) specific for R-LPS were used to localize the O-chain and R-LPS expressed in B. melitensis strain B115 by immunoelectron microscopy. Immunogold labelling was observed at the surface of B. melitensis B115 cells with the anti-R-LPS mAb but not with the anti-S-LPS mAb. In ultrathin sections, immunogold labelling with the S-LPS specific mAb was observed in the cytoplasm and in the periphery of the cytoplasm, probably at the cytoplasmic membrane. Immunogold labelling with the R-LPS specific mAb was observed at the outer membrane, outside the cells and on R-LPS vesicles. These results indicate that O-chain expressed in the rough B. melitensis strain B115 is immunogenic in mice, not exposed at the cell surface but present in the cytoplasm and most probably at the cytoplasmic membrane.

Introduction

The S-LPS of smooth-type Brucella spp. contains two distinct antigenic determinants designated A and M (Bundle et al., 1987, 1989; Diaz et al., 1968; Douglas & Palmer, 1988; Meikle et al., 1989; Wilson & Miles, 1932). The relative amounts of the two determinants vary among the smooth Brucella strains, and the determinants are absent on the rough strains, which lack O-chain (Alton et al., 1988). mAbs to S-LPS of Brucella spp. have been reported with specificity for the A-LPS determinant and the M-LPS determinant (Bundle et al., 1989; Douglas & Palmer, 1988; Palmer & Douglas, 1989). Other mAbs were found to react equally with S-LPS of both A- and M-dominant Brucella strains, some of them being specific to Brucella and others reacting also with Yersinia enterocolitica O:9 S-LPS (Douglas & Palmer, 1988; Palmer & Douglas, 1989). They were designated as specific for common (C) S-LPS determinants.

The rough Brucella melitensis strain B115, believed to be devoid of S-LPS or O-chain contamination, has been
used for preparing cytoplasmic antigens (Jones et al., 1973; Zygmunt et al., 1990) to study the immune response and also for extraction by the trichloroacetic acid method (Diaz et al., 1972, 1979, 1983; Fernandez-Lago et al., 1982; Moreno et al., 1981) of a polysaccharide fraction called polysaccharide B. Interest in Brucella polysaccharide B fraction stems in large part from its reported efficacy in tests that differentiate vaccinated from naturally infected cattle (Diaz et al., 1979, 1981, 1983). Recently, polysaccharide B fraction from smooth cells of B. melitensis 16M, produced by the trichloroacetic acid extraction method used for B. melitensis B115, was found to be composed of O polysaccharide (OPS) (up to 20% by weight) and cyclic D-glucan, serologically inactive, which was regarded as polysaccharide B (Bundle et al., 1987, 1988). In fact, polysaccharide B fraction from B. melitensis B115 rough cells was not really analysed. Nevertheless, Jones et al. (1973) showed that injection of living or killed cells of strain B115 into rabbits produced low-titre agglutinins to smooth cells, but they were unable to extract S-LPS from strain B115. Furthermore, whole cells of B. melitensis B115 were not able to absorb anti-smooth agglutinins from sera (Diaz et al., 1968).

We have reinvestigated the O-chain expression in B. melitensis B115 rough cells by using (1) hybridoma technology – characterization of one OPS-specific mAb induced by infecting mice with this rough strain, and (2) electron microscopic localization of the O-chain expressed in this strain by immunogold labelling using this O-polysaccharide-specific mAb.

**Methods**

**Bacterial strains.** The rough (R) Brucella melitensis strain B115 and the smooth (S) strains B. melitensis 16M and B. abortus 99 were obtained from the Institut National de la Recherche Agronomique, Laboratoire de Pathologie Infectieuse et Immunologie, Nouzilly, Tours, France. Purity and R and S phenotypes were clearly established by direct microscopic observation of colonies by obliquely reflected light and staining of R colonies with crystal violet (Alton et al., 1988). The R phenotype of B. melitensis strain B115 has been found stable over a period of 18 years (Alton et al., 1988; Jones et al., 1973).

Whole bacterial cells used for binding studies. Cultures were grown at 37°C for 48 h on trypticase soy agar (BioMérieux, Marcy-l’Etoile, France) slants supplemented with 0-1% (w/v) yeast extract (Difco). Cells were harvested by gentle agitation in sterile distilled water. Purity and phase (smooth, S; or rough, R) were checked using standard procedures (Alton et al., 1988). Dilutions were made in phosphate-buffered saline containing 0.5% (v/v) Tween 80 and the number of cells was determined by optical density measurements at 600 nm in a spectrophotometer (OD\textsubscript{600} = 0.165 for 10\textsuperscript{9} cells ml\textsuperscript{-1} for a 1 cm light path). Cells were killed by addition of a 3% (v/v) peracetic acid solution (Aldrich) (20 μl ml\textsuperscript{-1} to 10\textsuperscript{9} cells ml\textsuperscript{-1}) and incubated overnight at room temperature. Before use, killing was checked by taking 1 ml of each cell suspension and, after three washes in sterile distilled water, 0-2 ml was spread on Petri dishes containing the culture medium.

**Bacterial fractions.** Cell walls (CW) from B. melitensis B115 were isolated as previously described (Dubray & Charriaut, 1983). Briefly, cells were inactivated by heating at 65°C for 1 h and broken with glass beads in a Braun MSK homogenizer or a Dyno-Mill apparatus (W.A. Bachofen, Basel, Switzerland). Crude CW were recovered by centrifugation (at 53000 g, 4°C for 1 h). CW were obtained by treating crude CW with 1% (w/v) Triton X-100 in 0-2 M-NaCl, 0-01 M-MgCl\textsubscript{2} for 30 min at 20°C, washed six times in distilled water and lyophilized.

Cell extracts (CE) of B. melitensis B115 were obtained by ultrasonication. Cells were inactivated by heat at 65°C for 1 h, washed three times in 0.9% NaCl and ultrasonicated for 15 min in 1 mM-EDTA, 30 mM-Tris pH 8. The ultrasonicated cells were then centrifuged for 10 min at 4000 g, and the supernatant was recovered.

Cytoplasmic fraction (CP) of B. melitensis B115 was prepared as described by Zygmunt et al. (1990) by breaking heat-inactivated cells with glass beads in a Dyno-Mill apparatus followed by a first ultrafiltration with hollow-fibre carbides of 0.1 μm cut-off (Amicon). The ultrafiltrate was treated with ribonuclease and deoxyribonuclease (Worthington Diagnostics) followed by a second ultrafiltration with hollow-fibre carbides of 100 kDa cut-off (Amicon). The retentate was centrifuged at 50000 g for 16 h and the cleared supernatant (CP) was lyophilized.

**LPS fractions.** S-LPS fraction of B. melitensis 16M was prepared by the phenol/water method (Leong et al., 1970). R-LPS fraction of B. melitensis B115 was obtained by the phenol/water/chloroform/light petroleum method of Galanos et al. (1969). Proteinase-K-digested S-LPS fractions of B. melitensis 16M, B. abortus 99 and B. melitensis B115 were prepared as described previously (Garin-Bastuji et al., 1990). S-LPS fraction of B. melitensis B115 was isolated from CW. S-LPS fractions of B. melitensis 16M and B. abortus 99 were isolated from whole smooth bacterial cells. OPS of B. abortus 99 and B. melitensis 16M was extracted by autoclaving S cells in 2% (v/v) acetic acid/10% (w/v) NaCl as described by Jacques et al. (1991). The OPS was purified by high-performance liquid chromatography using a size-exclusion preparative TSK-G-2000-SW (600 × 25 mm) column (Zygmunt et al., 1988).

**mAbs.** The anti-S-LPS and anti-R-LPS mAbs were obtained as described previously (Cloeckaert et al., 1990). Briefly, BALB/c mice were infected by intraperitoneal (i.p.) injection of 10\textsuperscript{8} B. melitensis B115 rough cells. After 4 months, the mice were boosted i.p. with 10\textsuperscript{7} heat-Killed B. melitensis B115 rough cells. Four days after the booster injection, spleen cells were fused with cells of the NSO non-secreting myeloma cell line in a ratio of 5:1. After fusion, cells were suspended in selective hypoxanthine-aminopterin-thymidine-containing medium and seeded in 96-well microtitre plates at 5 x 10\textsuperscript{4} splenocytes per well. Anti-Brucella hybridomas were screened by ELISA and cloned by the limiting dilution technique. Specificity was determined by immunoblot analysis and ELISA using CW, CE and R-LPS of B. melitensis B115 and S-LPS of B. melitensis 16M and B. abortus 99. Ascitic fluids were prepared in BALB/c mice by the method of Limet et al. (1987).

**Reagents.** These were as follows. Glycine-buffered Saline (GBS): 0.17 M-NaCl, 0.1 M-glycine and 6.0 mM-Na\textsubscript{2}PO\textsubscript{4}; pH 9.2. GBS-EDTA-Tw: GBS containing 30 mM-EDTA and 0.1% Tween 80, final pH 9.2. Citrate-phosphate buffer: 0.051 M-Na, HPO\textsubscript{4}, 0.024 M-citric acid; pH 5.4. NaCl-Tw: 0.15 M-NaCl, 0.01% Tween 20. Tris-buffered saline (TBS): 0.15 M-NaCl, 10 mM-Tris/HCl; pH 7.5. Tw-TBS: TBS containing 0.05% (w/v) Tween 20. TBS-3% BSA: TBS containing 3% (w/v) bovine serum albumin. Tw-TBS-1% BSA: Tw-TBS containing 1% (w/v) BSA. PBS-Tw: phosphate-buffered saline (PBS): 0.15 M-NaCl, 0.01 M-sodium phosphate, pH 7.5) containing 0.05% Tween 20. PBS-1% BSA: PBS containing 1% (w/v) BSA.

**Antisera.** Rabbit anti-mouse immunoglobulin antiserum (RAM) was produced by repeated intradermal injections of 100 μg of mouse IgG.
After three injections at 15 d intervals, the rabbits were bled. Each month, boosters of 100 µg were repeated and followed by bleeding 10 d later. The best bleedings, as determined by latex agglutination immunoassay (Limet et al., 1988) using latex particles coated with mouse IgG, were pooled.

Rabbit anti-Brucella antiserum was produced by repeated intradermal injections of 10^8 killed cells of B. abortus 45/20 (R). After three injections at 15 d intervals, the rabbits were bled. The bleedings were tested by ELISA using B. abortus 45/20 (R) cells as antigen. Immunoglobulin fraction (Ig) was obtained by (NH_4)_2SO_4 precipitation.

** Peroxidase conjugation.** Protein A (Sigma) was conjugated with horseradish peroxidase (Sigma) using a modification of the method of Nakane & Kawanai (1974) as described by Dubray & Limet (1987). The protein/enzyme ratio was 1:2.

ELISA. Supernatants of hybridomas cultures or ascitic fluids were assayed for antibody activity by solid-phase ELISA against antigens coated by overnight incubation at 37°C after dilution in fivefold-diluted GBS in water (Voller et al., 1979). The antigens used to coat the plates consisted of CW (20 µg ml^-1), CE (diluted 1 in 1000), S-LPS (4 µg ml^-1), R-LPS (10 µg ml^-1), OPS (4 µg ml^-1) and whole bacterial cells (OD_550_1.5). Whole bacterial cells were immobilized on microtiter plates (Greiner-Labortechnik-Stuttgart) by means of rabbit anti-Brucella Ig adsorbed at a coating concentration of 10 µg ml^-1. Hybridoma supernatants were diluted (1 in 2) in GBS-EDTA-Tw. Binding of the antibodies to CW, CE, S-LPS or R-LPS was revealed by using RAM diluted 1 in 1000 in GBS-EDTA-Tw and peroxidase-conjugated protein A diluted 1 in 1000 in GBS-EDTA-Tw containing 2% (v/v) foetal calf serum. Binding to whole bacterial cells was revealed with peroxidase-conjugated goat anti-mouse immunoglobulins (Kierkegaard and Perry Laboratories) diluted 1 in 1000 in GBS-EDTA-Tw containing 2% foetal calf serum. Excess of reagents between the different incubations were removed by five washings with NaCl-Tw. o-Phenylenediamine (0.4%, w/v) and 2 M-H_2O_2 in citrate/phosphate buffer were used to reveal peroxidase activity. The ELISA titres were estimated as the highest dilution giving a difference in absorbance (492-620 nm) above twice the mean of the corresponding blank values.

** Thin-section preparation.** The method of thin-section preparation was as described by Lam et al. (1987). Samples were fixed in 1 ml of 2.5% (v/v) glutaraldehyde in PBS overnight at 4°C or in 1 ml 4% (v/v) paraformaldehyde buffered with 0.1 M-cacodylate at pH 7.25 for 1 h at room temperature. After fixation, samples were set in 5% (w/v) agarose and washed in buffer. The cores were then dehydrated through a stepwise alcohol series and infiltrated in propylene oxide. The samples were then embedded in Epon 812 resin and cured for 24 h at 37°C. Ultrathin sections were obtained by cutting the embedded material with an ultramicrotome (Reichert Jung Ultracut E) and deposited on 200-mesh gold grids.

** Immunogold labelling. (i) Whole-mount incubations.** A 5 µl sample of cell suspension in water (10^10 bacteria ml^-1) was deposited on carbon-Formvar-coated 200-mesh copper grids. After being air-dried (about 30 min), grids were incubated in the following reagents: PBS-3% BSA (30 min at 37°C), asartic fluid containing the mAb diluted 1 in 50 in PBS-Tw (2 h at 37°C), sheep anti-mouse biotinylated Ig (Amersham) diluted 1 in 200 in PBS-Tw (1 h at room temperature), gold-labelled streptavidin (15 nm) (Amersham) diluted 1 in 20 in PBS-Tw (1 h at room temperature). Washings between incubation periods were performed with NaCl-Tw. After four washings in NaCl-Tw and four washings in distilled water, grids were observed in a transmission electron microscope (Philips CM 10).

(ii) Section incubations. The same procedure as for whole mounts was followed, except that before incubation in PBS-3% BSA the grids were pretreated in 10% (v/v) H_2O_2 in distilled water for 30 min and then washed in distilled water. After immunogold labelling, the sections were stained with 4% (w/v) uranyl acetate in water and 0.4% lead citrate in 0.1 m-NaOH.

** Results**

** Monoclonal antibodies**

Twenty-two O-chain-specific mAbs were generated by fusion of spleen cells from mice infected with the rough B. melitensis strain B115 with the NSO myeloma. OPS specificity was determined by ELISA using OPS preparations of B. abortus 99 and B. melitensis 16M and by immunoblotting using proteinase-K-digested S-LPS preparations. One of these mAbs, designated 12G12, was subcloned and characterized by ELISA and immunoblotting to be further used for immuno-electron microscopy. According to its reactivity in ELISA with OPS and S-LPS of B. abortus 99 (A dominant) and B. melitensis 16M (M dominant) and the absence of reaction with R-LPS, mAb 12G12 recognized the OPS of both S-LPS types (Table 1). Immunoblotting using this mAb on proteinase-K-digested S-LPS (not shown), on S-LPS prepared by the phenol/water method and on an R-LPS preparation confirmed the ELISA results (see Fig. 4).

According to its reactivity with purified R-LPS in ELISA (Table 1) and immunoblotting (Fig. 4), mAb A68/03F03/D05 was specific for R-LPS. The weak reactivity of this mAb with S-LPS preparations in
Table 1. ELISA binding of the anti-S-LPS mAb I2G12 and the anti-R-LPS mAb A/6803F03/D05 to S-LPS and OPS from B. abortus 99 (A dominant) and B. melitensis 16M (M dominant), and to B. melitensis B115 bacteria and fractions

The values reported represent the absorbance value of a 1/2 hybridoma supernatant dilution after subtraction of the blank value.

<table>
<thead>
<tr>
<th>mAb</th>
<th>Isotype</th>
<th>S-LPS 99</th>
<th>S-LPS 16M</th>
<th>OPS 99</th>
<th>OPS 16M</th>
<th>ELISA against B. melitensis B115</th>
</tr>
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<tbody>
<tr>
<td>I2G12</td>
<td>IgG1</td>
<td>2.010</td>
<td>2.278</td>
<td>2.317</td>
<td>2.080</td>
<td>1.686 1.883 0.080 0.042</td>
</tr>
<tr>
<td>(S-LPS specific)</td>
<td></td>
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<td></td>
</tr>
<tr>
<td>A68/03F03/D05</td>
<td>IgG2b</td>
<td>1.150</td>
<td>0.623</td>
<td>0.017</td>
<td>0.259</td>
<td>1.409 1.864 1.865 0.992</td>
</tr>
<tr>
<td>(R-LPS specific)</td>
<td></td>
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ELISA (Table 1) was due to the presence of R-LPS molecules in the preparations (as also observed in Fig. 4).

**ELISA**

Binding of anti-S-LPS mAb I2G12 to CE, CW, R-LPS and whole bacteria of B. melitensis B115 was compared with the binding of the anti-R-LPS mAb A68/03F03/D05 (Table 1). Binding to CE and CW was observed for both mAbs. Binding to R-LPS was only observed for the R-LPS-specific mAb. In contrast to the R-LPS-specific mAb, the S-LPS-specific mAb did not bind to whole bacteria of B. melitensis B115, which confirms the rough phenotype of this strain. These results indicate that O-chain is present in this rough strain but that it is not exposed on the cell surface.

**Whole-mount immunogold labelling**

Immunogold labelling of whole B. melitensis B115 bacteria confirmed the absence of O-chain exposure on the cell surface: immunogold labelling was observed with the R-LPS-specific mAb A68/03F03/D05 but not with the S-LPS-specific mAb I2G12 (Fig. 1).

**Section immunogold labelling**

Immunogold labelling of sections of B. melitensis B115 cells using the S-LPS-specific mAb I2G12 revealed the presence of O-chain in the cytoplasm and at the periphery of the cytoplasm, probably at the cytoplasmic membrane (Fig. 2). No immunogold labelling was observed at the outer membrane by using this mAb. In some cells labelling was observed as clusters in the cytoplasm close to the cytoplasmic membrane (Fig. 2b), but in most of the cells immunogold labelling was dispersed in the cytoplasm. Immunogold labelling using the anti-R-LPS mAb A68/03F03/D05 was observed in the cytoplasm, near the cytoplasmic membrane and at the outer membrane, and outside the cells on R-LPS vesicles (Fig. 2c). No immunogold labelling was observed with a negative control mAb (Fig. 2d).

**Immunoblotting of B. melitensis B115 LPS**

B. melitensis B115 CW, CE, CP and R-LPS fractions were analysed by immunoblotting after SDS-PAGE (Fig. 4). The S-LPS-specific mAb I2G12 revealed bands in CE, CW and CP ranging from 30 to 43 kDa which were not revealed by the R-LPS specific mAb A68/03F03/D05. The latter mAb revealed bands of lower molecular mass, below 21 kDa. The S-LPS-specific mAb I2G12 did not react with the B. melitensis B115 R-LPS fraction (lane 4) extracted according to the procedure of Galanos et al. (1969), which corresponds to R-LPS of the cell surface.

Proteinase-K-digested S-LPS fraction from B. melitensis B115 CW showed the same molecular mass as
proteinase-K digested S-LPS fraction from *B. abortus* 99 and *B. melitensis* 16M (not shown).

**Discussion**

The present study reports the O-chain expression and its localization in the *B. melitensis* B115 rough strain which is commonly used for cytoplasmic antigen extraction (Jones *et al.*, 1973; Zygmunt *et al.*, 1990) and for polysaccharide B extraction (Diaz *et al.*, 1972, 1983, 1979; Moreno *et al.*, 1981). The interest of using *B. melitensis* B115 was that it was believed to be devoid of S-LPS or O-chain since this strain is considered as a rough strain by classical criteria (Alton *et al.*, 1988).

We obtained 22 OPS-specific mAbs by fusion of spleen cells of mice infected with *B. melitensis* strain B115 with the NSO myeloma, suggesting expression of immunogenically active O-chain in this strain. Most of the mAbs recognized epitopes common to both A and M S-LPS types (data not shown). mAbs to S-LPS of *Brucella* spp. have already been reported with specificity for common epitopes and cross-reacting or not with *Y. enterocolitica* O:9 S-LPS (Palmer & Douglas, 1989).

As shown by ELISA the anti-S-LPS mAb 12G12 did not bind to whole *B. melitensis* B115 cells but it bound well to CE or CW of this strain. The mAb specific for R-LPS bound to whole bacterial cells as well as to CE or CW. Thus this rough strain expresses O-chain but the latter is not exposed at the cell surface. This is also confirmed by the fact that mAb 12G12 did not react with the *B. melitensis* B115 R-LPS fraction prepared from whole bacterial cells by the phenol/water/chloroform/light petroleum method of Galanos *et al.* (1969). This method leaves the bacterial cells morphologically intact as shown by electron microscopy (Dubray, 1981). Therefore R-LPS extracted by this method is probably only the cell-surface-exposed R-LPS; this may explain why Jones *et al.* (1973) were unable to extract S-LPS from *B. melitensis* B115. Thus, rabbits immunized with R-LPS prepared from *B. melitensis* B115 by the method of Galanos *et al.* (1969) produced agglutinins to rough brucellae only and precipitins to the R surface antigen only (Jones *et al.*, 1973).

O-chain expression could be visualized by immunoelectron microscopy in thin sections of cells of *B. melitensis* B115. The localization of O-chain in this strain is mainly cytoplasmic. No immunogold labelling of the outer membrane was observed by using the S-LPS
Fig. 2. Immunogold labelling of sections of *B. melitensis* B115 cells with anti-S-LPS mAb 12G12 (*a*, *b*), anti-R-LPS mAb A68/03F03/D05 (*c*) and negative control mAb (*d*). Bars, 0.5 μm.
Fig. 3. Immunogold labelling of sections of *B. melitensis* B115 CW with anti-S-LPS mAb 12G12 (a) and negative control mAb (b). Bars, 0.5 μm.

Fig. 4. Immunoblot of *B. melitensis* B115 CE (125 μg) (lanes 1 and 6), CW (125 μg) (lanes 2 and 7), CP (125 μg) (lanes 3 and 8), R-LPS (62.5 μg) (lanes 4 and 9) and *B. melitensis* 16M S-LPS (phenol fraction) (62.5 μg) (lanes 5 and 10) with anti-S-LPS mAb 12G12 (lanes 1, 2, 3, 4, 5) and anti-R-LPS mAb A68/03/F03/D05 (lanes 6, 7, 8, 9, 10). The positions of protein molecular mass markers are shown on the left.
specific mAb. In contrast the anti-R-LPS mAb, a marker of the outer membrane, also bound to whole bacterial cells.

We do not know yet if the O-chain expressed in the cytoplasm is free or linked to R-LPS or to the lipid carrier. The transfer of the OPS to the core is a complex reaction between membrane-bound molecules, the undecaprenol-diphosphate-polysaccharide on the one hand, and the complete core on the other (Måkelä & Stocker, 1984; Mulford & Osborn, 1983; Osborn, 1979). Since S-LPS biosynthesis takes place in the cytoplasmic membrane, a part of the O-chain expressed in the rough B. melitensis B115 strain is probably linked to a lipid moiety: R-LPS or the undecaprenol-diphosphate-polysaccharide lipid carrier. Several hypotheses can be put forward to explain the rough phenotype of B. melitensis B115 despite its O-chain expression: the core is incomplete, as in Salmonella rfaK mutants (Beckmann et al., 1964; Kent & Osborn, 1968; Lindberg et al., 1972; Weiner et al., 1965); unfinished O units are not attached to the core in vivo (Yuasa et al., 1969); or, if both requirements are fulfilled (1) transfer of the O-chain from the undecaprenol-diphosphate-polysaccharide carrier to the core does not occur, or (2) translocation of S-LPS molecules from the cytoplasmic membrane to the outer membrane does not occur.

The lipid-bound O-chain has been isolated from B. melitensis B115 CW as evidenced by SDS-PAGE and immunoblot analysis. Bands revealed by the anti-S-LPS mAb 12G12 have the same molecular size as S-LPS bands of B. melitensis 16M and B. abortus 99 smooth strains. To clearly identify the reason for the absence of S-LPS at the cell surface, high-molecular-mass proteinase-K-digested LPS from CW should be purified, acid hydrolysed and the lipid moiety analysed as well as the polysaccharide.

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