Immunochemical Characterization of a Polysaccharide Antigen of *Bacteroides fragilis* with an IgM Monoclonal Antibody

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An IgM mouse monoclonal antibody (McAb) Bf4 was produced to a surface polysaccharide of *Bacteroides fragilis* NCTC 9343. Immunoblotting showed that McAb Bf4 reacted strongly with a high molecular mass structure which was sensitive to oxidation with periodate but resisted protease treatment. An inhibition enzyme-linked immunosorbent assay (ELISA) indicated that McAb Bf4 did not cross react with the sixteen *Bacteroides* species and strains tested. Cells of *B. fragilis* NCTC 9343 recovered from the various interfaces of a Percoll discontinuous density gradient were tested in the inhibition ELISA. Bacteria from the 0–20%, 20–40% and 40–60% interfaces inhibited the ELISA; however, cells from the 60–80% interface did not. Electron microscopy with immunogold labelling showed that McAb Bf4 did not react with the extracellular fibrous network on bacteria recovered from the 0–20% interface, or the extracellular electron dense layer on cells from the 60–80% interface; however, it was associated with a surface structure on cells from the 20–40% interface. Growth *in vivo* did not enrich for bacteria with this structure.

INTRODUCTION

*Bacteroides fragilis* is the most common Gram-negative anaerobic organism isolated from clinical specimens (Gorbach & Bartlett, 1974; Duerden, 1980). However, the factors responsible for the enhanced virulence of this species are unknown despite the development of a number of models for studying the pathogenic mechanisms of bacteroides (Hofstad, 1984). Previous reports have suggested that the capsular polysaccharide of *B. fragilis* may represent a virulence factor (Onderdonk et al., 1977; Connolly et al., 1984).

A number of *B. fragilis* surface antigens have been identified (Kasper, 1976; Babb & Cummins, 1978; Cousland & Poxton, 1983; Lambe et al., 1984; Weintraub et al., 1985), but little is known about the distribution and expression of these structures. Variation in the degree of capsulation of *B. fragilis* occurs during passage *in vitro* and *in vivo* (Kasper et al., 1980; Patrick & Reid, 1983; Patrick et al., 1984). Cells of *B. fragilis* with different sizes of capsule can be separated on a Percoll density gradient and appear to differ structurally when examined by electron microscopy (Patrick et al., 1986). The relationship between the surface structures observed by us and either the surface polysaccharide described as 'capsule' by Kasper (1976) or the glyocalyx reported by Lambe et al. (1984) is unknown. Characterization of these antigens using highly specific monoclonal antibodies (McAbs) and immunochemical techniques, e.g. immunoblotting and electron microscopy with immunogold labelling, will provide a way of comparing the surface structures identified by different research groups. McAbs have been produced to rough LPS molecules of *B. fragilis* (Linko-Kettunen et al., 1984) and a high molecular mass surface structure on *B. fragilis* (Reid et al., 1985).

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Abbreviations: McAb, monoclonal antibody; TBS, Tris-buffered saline; CB, cacodylate/HCl buffer; PCP, phenol–chloroform–petroleum.

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In this study immunoblotting was used to define the immunochemical nature of the structure recognized by McAb BF4 and an inhibition enzyme-linked immunosorbent assay (ELISA) was used to determine the specificity for Bacteroides strains and species. The distribution of the BF4 epitope on B. fragilis populations grown in vitro and in vivo was investigated using immunofluorescence and electron microscopy with immunogold labelling.

**METHODS**

*Bacterial growth conditions.* The Bacteroides strains used in this study were as follows: B. fragilis NCTC 9343 from the Department of Microbiology and Immunobiology, the Queen's University of Belfast, UK; B. fragilis GNAB from the Department of Bacteriology, Edinburgh University Medical School, UK; B. ovatus ATCC 8483, B. distasonis ATCC 8503, B. thetaiotaomicron NCTC 10582, B. fragilis ATCC 23745; NCTC 9344, NCTC 10584, NCTC 8560 and clinical B. fragilis isolates Bf12, Bf17, Bf18, Bf19, Bf20, Bf21, Bf25 and Bf30 from the Department of Medical Microbiology, St Bartholomew's Hospital Medical College, London, UK. B. fragilis strains grown to late exponential phase in defined broth (van Tassell & Wilkins, 1978) or on horse blood agar plates were incubated at 37°C in an anaerobic chamber (Forma) with an atmosphere of 80% (v/v) N2, 10% (v/v) H2 and 10% (v/v) CO2. Identification was confirmed using the API system.

Production of McAbs. A BALB/c mouse was immunized with whole cells of B. fragilis NCTC 9343. The mouse was inoculated intraperitoneally each week for 3 weeks with 0.2 ml of a bacterial suspension of 1-0 × 10^8 c.f.u. in one-quarter strength Ringer's solution, and an additional inoculation of 0.2 ml was given 4 d before fusion. Spleen cells from the intraperitoneally moused were fused with x 63 Ag-8.653 myeloma cells using the polyethylene glycol fusion technique (Lemke et al., 1978). Hybrid cell lines were selected in RPMI 1640 medium containing hypoxanthine–aminopterin–thymidine and 15% (v/v) foetal calf serum (Gibco). Hybridoma culture supernate was screened for antibody to B. fragilis NCTC 9343 by ELISA (see below). Hybridoma culture supernate was used in all experiments.

The hybridoma cell line BF4 was cloned using a limiting dilution method and BALB/c macrophages were used to assist the growth of hybridoma clones. Cells were concentrated by centrifugation and 1 ml samples stored at −80°C in growth medium containing 15% (v/v) foetal calf serum and 10% (v/v) dimethyl sulphoxide. Hybridoma cells from this stock were used to produce large quantities of hybridoma BF4 culture supernate.

**ELISA.** An indirect ELISA was used to screen for McAb producing hybridoma cell lines. Wells of PVC microtitre plates (Becton and Dickenson) were each coated with bacteria (1 × 10^6 cells ml−1) suspended in 50 mM-sodium carbonate buffer, pH 9.6. The plates were incubated at 37°C for 2 h and washed three times with 50 mM-sodium phosphate buffer, pH 7.4, containing 150 mM-NaCl (PBS) and 0.05% (v/v) Tween 20. Plates were stored at −20°C.

Hybridoma culture supernate (100 μl) was added to each well, and the plates were incubated at 37°C for 1 h and washed as before [dilutions, if required were made in PBS containing 1% (w/v) bovine serum albumin (BSA–PBS)]. Goat anti-mouse IgM (Mu chain specific) and IgG-alkaline phosphatase conjugates (Tago) diluted 1 in 3000 in 1% (w/v) BSA–PBS were added to each well (100 μl) and the plates were incubated at 37°C for 1 h. After washing, 100 μl of p-nitrophenyl phosphate (Sigma) (1 mg ml−1) in 50 mM-sodium carbonate buffer, pH 9.8, containing 1 mM-MgCl2 was added to each well. The plate was incubated at 37°C for 1 h; the reaction was stopped with 100 μl 3 M-NaOH and absorbance (at 405 nm) was read using a BioRad automatic plate reader.

An inhibition ELISA was used to test for cross reactivity with other bacteroides. Bacteria suspended in 0.5 ml 2% (w/v) BSA in PBS to a concentration of 5.0 × 10^6 c.f.u. ml−1 were mixed with 0.5 ml of hybridoma culture supernate containing McAb BF4 and incubated at 4°C for 18 h. After centrifugation at 5000 g, triplicate samples (100 μl) of the supernate were tested on ELISA plates coated with B. fragilis NCTC 9343.

An ELISA was used to determine the immunoglobulin class of McAb BF4. Sheep anti-mouse IgG2a (Serotec), goat anti-mouse IgG1 (Sigma), IgG2b (Sigma), IgG3 (Sigma) and IgM (Sigma) were diluted in 50 mM-sodium carbonate buffer, pH 9.6, and coated onto PVC microtitre plates as before. Hybridoma culture supernate (50 μl) was added to each well; the plates were incubated for 2 h at 37°C, and then washed as before. Goat anti-mouse IgG (specific for light and heavy chains) conjugated to horse-radish peroxidase (Sigma) was diluted 1 in 1000 in 1% (w/v) BSA in PBS and 50 μl was added to each well. The plates were incubated at 37°C for 1 h and washed as above. Orthophenylene diamine (Sigma) substrate (40 mg) was dissolved in 100 ml of a solution containing 100 mM-citric acid, 200 mM-Na2HPO4 and 0-01% (w/v) H2O2; 50 μl of the solution was added to each well and the plates were incubated for 30 min; the reaction was stopped with 1 ml H2SO4 and the absorbance read at 520 nm.

*Enrichment for bacterial variants.* Capsulate and non-capsulate B. fragilis (defined by light microscopy) were separated by Percoll density gradient centrifugation (Patrick & Reid, 1983).

*Preparation of antigens.* Antigens were extracted from bacteria by the aqueous phenol method of Westphal & Ludertz (1954). The aqueous phase was centrifuged at 10000 g for 30 min at 4°C to remove insoluble material and the supernate was stored at −20°C.

Antigens from bacteria were also extracted with a glycine buffer containing EDTA. Bacteria harvested from
500 ml broth were resuspended in 10 ml 3 mM-glycine buffer, pH 5-0, containing 10 mM-EDTA, and were incubated at 37 °C for 30 min. Whole cells were removed by centrifugation at 10000 g for 30 min at 4 °C and the extract was stored at −20 °C. Extracts were treated with protease or sodium periodate by the method of Cousland & Poxtin (1984).

**Carbohydrate assay.** The anthrone method of Morris (1948), with d-glucose as a standard, was used.

**SDS-PAGE.** This was done on 10% vertical slab gels using the Laemmli buffer system (Laemmli, 1970) at a constant current of 40 mA per gel.

**Immunoblotting and enzyme immunoassay.** The method of Tewbin et al. (1979) was used with the following modifications. Material was transferred to a nitrocellulose membrane (Transblot TM transfer medium; BioRad) in 25 mM-Tris/192 mM-glycine buffer, pH 8.3, containing 20% (v/v) methanol at 350 mA for 1.5 h. After washing for 10 min in Tris-buffered saline (TBS; 20 mM-Tris/HCl, 500 mM-NaCl, pH 7.5) the membrane strip containing the protein molecular mass standards was fixed with 1% (v/v) acetic acid and stained with amido black. The rest of the membrane was placed in 3%(w/v) BSA in TBS (BSA-TBS) for 16 h at 4 °C. It was then transferred into hybridoma culture supernate, or an appropriately diluted mouse derived antisera, and incubated for 2 h at room temperature. After three washes in 0.025% Tween 20 in TBS, the membrane was placed in goat anti-mouse IgM-alkaline phosphatase conjugate (Tago) specific for the Mu chain, then diluted 1 in 3000 in 1% (w/v) BSA in TBS and incubated for 1 h at room temperature. The membrane was washed three times in TBS and placed in 50 mM-Tris/HCl buffer, pH 8.2, containing Fast Red (3 mg ml−1) and naphthol-AS-MX-phosphate (0.2 mg ml−1). Colour development took place in 5–15 min, and the reaction was then stopped by washing in distilled water. All the above steps were done with gentle agitation throughout.

**Immunogold labelling and electron microscopy.** Bacteria were washed twice in 100 mM-sodium cacodylate/HCl buffer, pH 7.2 (CB), and fixed in CB containing 2% (v/v) paraformaldehyde and 0.1% (v/v) glutaraldehyde for 1 h at 4 °C. The cells were again washed in CB, dehydrated in graded alcohols and embedded in LR white resin. For gold labelling, ultrathin sections on nickel grids were treated with McAb Bf4 followed by goat anti-mouse IgM-alkaline phosphatase conjugate (Tago) specific for the Mu chain, then diluted 1 in 3000 in 1% (w/v) BSA in TBS and incubated for 1 h at room temperature. The membrane was washed three times in TBS and placed in 50 mM-Tris/HCl buffer, pH 8.2, containing Fast Red (3 mg ml−1) and naphthol-AS-MX-phosphate (0.2 mg ml−1). Colour development took place in 5–15 min, and the reaction was then stopped by washing in distilled water. All the above steps were done with gentle agitation throughout.

**RESULTS**

**SDS-PAGE and immunoblotting of the phenol–water extract**

Reactions with polyclonal mouse antiserum. SDS-PAGE and immunoblotting with homologous anti-\(B.\) fragilis NCTC 9343 antiserum detected multiple bands (Fig. 1c). A diffuse staining pattern and a series of closely spaced discrete bands (arrowed S) were detected at the top of the blot. A series of regularly spaced discrete bands was observed in the middle of the blot (labelled R + n). At least two heavily stained bands were seen in the lower region of the blot.

Reaction with McAb Bf4. Staining was observed only in the high molecular mass region when McAb Bf4 was used for immunoblotting (Fig. 1b). McAb Bf4 reacted strongly with both the diffuse staining region and the closely spaced discrete bands.

Treatment of antigen with protease, periodate and heat. The structure recognized by McAb Bf4 resisted protease digestion (result not shown) and the immunoblot was identical to that for the control antigen (Fig. 1b). Oxidation of the antigen preparation with sodium periodate destroyed the Bf4 antigenic determinant (Fig. 1a). Heating at 100 °C for 3 min did not destroy the structure containing the Bf4 epitope.

**SDS-PAGE and immunoblotting of the glycine–EDTA extract**

Immunoblotting of the glycine–EDTA extract after SDS-PAGE with McAb Bf4 revealed multiple closely spaced discrete bands and at least four extra bands in the high molecular mass region of the blot (Fig. 2b, d). Protease treatment did not affect the closely stained bands but the extra bands in the top region of the blot disappeared (Fig. 2c). Oxidation with sodium periodate removed the closely spaced bands, but multiple faintly immunoreactive bands were detected in the lower region of the blot after treatment (Fig. 2a).

**ELISA**

McAb Bf4 was identified as an IgM antibody using a sandwich ELISA. Inhibition studies were done to test the reactivity of McAb Bf4 with other \(Bacteroides\) species.
and strains. Also, cells of *B. fragilis* NCTC 9343 recovered from the various interfaces of a Percoll discontinuous density gradient were tested for expression of the Bf4 antigen.

*B. ovatus* ATCC 8483, *B. distasonis* ATCC 8503 and *B. thetaiotaomicron* NCTC 10582 did not inhibit the ELISA. Four reference strains and nine clinical isolates of *B. fragilis* were also tested. Inhibition was detected only with the homologous strain of NCTC 9343.

Bacteria recovered from the 0–20%, 20–40% and 40–60% interfaces of a Percoll discontinuous density gradient inhibited the ELISA, but cells from the 60–80% interface did not. No attempt was made to measure the minimum number of cells required to inhibit the assay.

**Immunofluorescence and immuno-electron microscopy**

Immunofluorescence and immuno-electron microscopy of cells taken from the four interfaces of the Percoll gradient gave the same pattern of labelling. The absence of the Bf4 epitope from the 60–80% interface was confirmed. Cells from this interface are apparently non-capsulate by light microscopy, but have an extracellular electron dense layer by electron microscopy (Patrick...
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Fig. 2. SDS-PAGE and immunoblot analysis of antigens extracted from B. fragilis NCTC 9343 with mild heat and EDTA-glycine buffer, pH 5.0. Antigen treatments were periodate (a) with control (b), and protease (c) with control (d). Carbohydrate loaded was 50 µg per lane.

et al., 1986). A low proportion of cells from the 0–20% (Fig. 3a) and 40–60% (not illustrated) interfaces were labelled, but cells from the 20–40% interface were enriched for the Bf4 epitope (Fig. 3b). The surface structures of B. fragilis are not easily resolved with the fixation procedures required to preserve the reactivity of the antigens for immunoelectron microscopy; therefore duplicate samples were stained with ruthenium red. The results indicate that the Bf4 epitope is associated with the marginal fibrous network present on cells from the 20–40% interface (Fig. 3d), but not with the extensive fibrous network which is present on cells from the 0–20% interface (Fig. 3c). Two other monoclonal antibodies (Bf1; Reid et al., 1985 and Bf2; not illustrated) give a similar labelling pattern for each of the interfaces and could not be related to a specific surface structure.

Cells from the 0–20% interface were grown for 24 h in chambers implanted in the mouse peritoneal cavity (Patrick et al., 1984). These populations were not enriched for the Bf4 epitope (Fig. 4). Similar labelling patterns were seen with these populations before and after in vivo passage (Figs 3a and 4).

DISCUSSION

Immunoblotting and immunoelectron microscopy showed that McAb Bf4 reacted with an epitope on a high molecular mass surface polysaccharide. This structure was distinct from the
Fig. 3. Electron micrographs of cells of *B. fragilis* NCTC 9343 recovered from the interfaces of a Percoll discontinuous density gradient. (a) Cells from the 0–20% interface labelled with McAb Bf4 and anti-mouse IgM immunogold conjugate; note the low proportion of gold labelled cells. (b) Cells from the 20–40% interface labelled as in (a); note that most of the cells are labelled. (c) Cells from the 0–20% interface stained with ruthenium red; note the extensive fibrous network (*). (d) Cells from the 20–40% interface stained with ruthenium red; note the marginal fibrous network (F) adjacent to the outer membrane (OM); CM, cytoplasmic membrane. Bar = 0.5 μm in main micrographs and 100 nm in Fig. 3(d) insert.
extensive fibrous network, present on cells recovered from the 0–20% interface of a Percoll density gradient, and the electron dense layer, present on cells from the 60–80% interface. It was, however, associated with the marginal fibrous network observed on cells from the 20–40% interface. These results indicate that structures which look different by electron microscopy may also be antigenically different. Immunogold labelling of cells grown in the mouse peritoneal cavity gave a heterogeneous labelling pattern which indicates that this epitope was not selected \textit{in vivo}.

ELISA inhibition studies showed that McAb Bf4 recognized a type-specific surface antigen. Kasper \textit{et al.} (1983) showed that extracted capsular polysaccharide from \textit{B. fragilis} NCTC 9343 was chemically and antigenically different from that of \textit{B. fragilis} ATCC 23745, and type-specific epitopes were identified in the capsular material. This capsular material was detected in the high molecular mass region of a 20\%(w/v) acrylamide gel by silver staining (Weintraub \textit{et al.}, 1985). Weintraub \textit{et al.} (1985) also extracted LPS from \textit{B. fragilis} NCTC 9343 by the hot phenol–water method and subsequently extracted the aqueous phase with a phenol–chloroform–petroleum (PCP) mixture: rough LPS only was detected with SDS-PAGE and silver staining. However, \textit{B. fragilis} NCTC 9343 could also contain smooth LPS with lipopolysaccharide side chains since the PCP extraction procedure preferentially extracts lipophilic rough LPS while smooth LPS is excluded (Galanos \textit{et al.}, 1969). Cousland & Poxton (1983) have observed heterogeneity in LPS extracted from a variety of \textit{B. fragilis} strains by the hot phenol–water method. They separated LPS extracted from \textit{B. fragilis} GNAB 92 by electrophoresis on a 10\%(w/v) acrylamide gel and detected possible smooth LPS (labelled S) in the top of the blot.
McAb Bf4 reacted with B. fragilis NCTC 9343 antigens in this region of the blot. McAb Bf4 did not react with the typical LPS 'ladder' (Fig. 1b, labelled R + n) in the middle of the blot, suggesting that either the Bf4 epitope was not part of a repeating O-side-chain or the concentration of the LPS containing short O-side-chains was below a detectable level when using McAb Bf4. The epitope could be associated with a repeating capsular polysaccharide unit but the polymer would need to be covalently linked to a lipid to bind SDS, or have a negative charge for electrophoresis to occur.

E. coli LPS formed non-covalently linked multimers during SDS-PAGE when 0.1% (w/v) SDS was used in 15% (w/v) acrylamide separating gels (Peterson & McGroarty, 1985). The multimers dissociated when 0.5% (w/v) SDS was used or when separating gels were overlayed with butanol and left overnight. McAb Bf4 could also be reacting with multimers of B. fragilis NCTC 9343 LPS. The SDS-PAGE conditions were not altered in this study but the hypothesis could be tested by increasing the SDS concentrations in both the sample buffer and the separating gel, or by electro-eluting the high molecular mass material from the top section of the gel, and re-analysing the material by SDS-PAGE and immunoblotting with McAb Bf4.

Poxton et al. (1985) detected protein–LPS complexes when outer membrane extracts of Pseudomonas aeruginosa were immunoblotted with McAbs specific for the LPS O-antigen. This report describes a similar observation using a glycine–EDTA extract of B. fragilis NCTC 9343 and immunoblotting with McAb Bf4. There are a number of possible explanations for this phenomenon. First, protein–LPS complexes released by the glycine–EDTA treatment could resist solubilization with SDS. Second, antigens could be completely solubilized by detergent but co-migrate to the same position in the gel. Third, solubilized high molecular mass peptides could become entangled with long polysaccharide chains containing the Bf4 epitope during electrophoresis. The stability of antigen complexes could be investigated by affinity chromatography. The structure could then be chemically analysed to determine whether it is a polysaccharide or a lipopolysaccharide.

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