Identification of an endoflagellar associated protein in Borrelia burgdorferi

H. EIFFERT*, T. SCHLOTT, M. HOPPERT†, HANNELORE LOTTER and R. THOMSSEN

Department of Medical Microbiology and †Institute of Microbiology, University of Göttingen, Germany

Summary. DNA of Borrelia burgdorferi was cleaved by the endonuclease EcoRI and ligated with the bacteriophage expression vector λgt11. After infection of the Escherichia coli strain Y1089, the plaques of recombinant phages were screened with a B. burgdorferi antiserum (human) for fusion proteins containing borrelia antigens. A positive clone produced a hybrid protein (p200) of c. 200 Kda. The corresponding native borrelia protein (p97) was identified as having an M, of 97 Kda. To localise protein p97 in the B. burgdorferi cell, immuno-electronmicroscopy and a Western blot of isolated flagella were used. Antibodies directed against proteins p200 and p97 recognised epitopes associated with the flagella.

Introduction

Borrelia burgdorferi is the causative agent of Lyme disease, a multisystem disorder characterised by various clinical manifestations at different stages. The bacterium represents a new species of Borrelia and is related to other spirochaetal pathogens such as Treponema and Leptospira. Little is known about the protective role of antibodies or about pathogenesis. Some of the major antigens of B. burgdorferi have been investigated by gel electrophoresis and immunological methods, and partly by cloning and nucleotide sequencing. The best characterised molecules are the outer-surface proteins OspA, OspB and pC,4-8 the major flagellar protein p419-" and the "common antigen" with an M, of c. 60 Kda.12 The common antigen and p41 share epitopes with other bacteria. In this report a further protein of B. burgdorferi is described.

Materials and methods

Bacterial strains

B. burgdorferi strain IRS Us was isolated, as described by others, from a tick (Ixodes ricinus) found near Göttingen. E. coli strains JM 109, Y1089 and Y1090 were provided by Stratagene Ltd, Heidelberg, Germany.

Cultivation of B. burgdorferi and antigen preparation

Large scale cultivation was by the method of Barbour. The strain was grown in a modified BSK medium for 5 days at 33°C. The spirochaetes were harvested by centrifugation at 10 000 g for 30 min at 4°C. The pellet was washed three times in phosphate-buffered saline (PBS) pH 7.4 and resuspended. The suspension was disintegrated in an ice bath by means of three 5-s bursts from a sonifier (Branson Sonic Power Co., Danbury, USA) operating at 30% maximal intensity. The preparation was then centrifuged at 10 000 g for 30 min at 4°C. The supernate containing borrelia antigen was stored in small volumes at -24°C.

Serum samples

These were obtained from the B. burgdorferi diagnostic laboratory, Department of Medical Microbiology, University of Göttingen, Germany. The serum used for screening for the fusion proteins was obtained from a forestry worker with clinical symptoms of Lyme disease and a high concentration of B. burgdorferi antibodies. The antibodies were detected by an IgM capture ELISA and an IgG ELISA.

Preparation of B. burgdorferi DNA

A culture (500 ml) was prepared by a modification of a method described previously. After centrifugation, the cells were suspended in 1 ml of 50 mM Tris HCl, pH 7.6, containing sucrose 25% w/w and lysozyme (Boehringer, Mannheim, Germany) 2 mg and incubated for 30 min at 37°C. Thereafter, 0.4 ml of 0.25 M edetic acid, pH 8.0, and 0.16 ml of sodium dodecylsulphate (SDS) 10% w/w were added and mixed. This lysate was digested with 0.5 mg of proteinase K (Boehringer) for 60 min at 37°C and extracted with phenol-chloroform; the DNA was then precipitated with ethanol.

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* Correspondence should be sent to Dr H. Eiffert, Hygiene Institut, Kreuzbergring 57, D-3400 Göttingen, Germany.
Preparation and screening of the expression library

The isolated DNA of B. burgdorferi was digested completely by the restriction enzyme EcoRI (Pharmacia, Uppsala, Sweden). For removal of RNA, DNase-free RNAase 0.1 mg/ml was added and incubated at 37°C for 60 min. The process of cleavage was monitored by agarose gel electrophoresis. After extraction with phenol-chloroform and precipitation with ethanol, the DNA was dried in a vacuum.

The DNA was ligated into EcoRI-digested, dephosphorylated λgt11 DNA (Bethesda Research Laboratories, Gaithersburg, USA) and packaged with DNA Packaging Kit (Boehringer) as described by the manufacturers. A suspension (100 µl) of the competent E. coli strain Y1089 was transfected with 1 µl of the suspension of recombinant bacteriophages for 60 min at 37°C and then added to 3 ml of soft agar containing 10 mM MgSO₄, ampicillin 0.01% and 0.3 mM isopropyl-β-D-thiogalactoside (IPTG). The soft agar was then poured on to Luria broth (LB) plates containing ampicillin 0.01% and incubated overnight at 37°C. Replicas of the plates were obtained by carefully placing dry nitrocellulose filter disks (Schleicher and Schüll, Dassel, Germany) on the soft agar for 30 min at 37°C. These were then removed and the position was marked on the plate with a needle. The filters were rinsed with PBS and incubated with milk powder (Uelzena, Uelzen, Germany) 5% w/w in 5 ml of PBS for 30 min at room temperature. Human sera with B. burgdorferi antibodies, or negative human sera, were then added in a final dilution of 1 in 1000 and incubated for 60 min at room temperature. After four 5-min washes with Tween 0.05% w/w in PBS, positive clones were detected by standard methods with peroxidase labelled rabbit antibodies against NaClO₄. The proteins were separated by a standard SDS-polyacrylamide gel electrophoresis (PAGE). The gels were stained with Coomassie Blue. Electrobetting was performed with a polyvinylidene difluoride membrane (PVDF, Millipore, Bedford, USA) as described by Gültken and Heermann. For the detection of bound antibodies, peroxidase labelled anti-human or anti-rabbit antibodies (Dako) were used.

Elution of specific antibodies

For the elution of specific antibodies from blot membranes, the technique described by Olmsted was used.

Electro-elution

Fusion protein was electro-eluted from a preparative SDS-polyacrylamide gel into a membrane trap (Bio-trap, Schleicher und Schüll).

Absorption procedure

E. coli lysate (100 µg), before or after infection with the recombinant bacteriophage, or electro-eluted protein p200 (30 µg) was mixed with 100 µl of serum containing B. burgdorferi antibodies. Absorbed and untreated control samples were held at room temperature for 4 h.

Immunisation of the rabbits

Rabbits were immunised subcutaneously with 0.8 mg of electro-eluted protein p200 after dialysis against NaCl 0.9% w/v. To 1 ml of the solution, 1 ml of Freund’s complete adjuvant was added. The immunisation procedure was repeated three times at intervals of 2 weeks. The negative control serum was obtained before immunisation. The specific antiserum used for immuno-electronmicroscopy was purified by affinity chromatography on protein A-Sepharose CL-4B (Pharmacia, Uppsala, Sweden).

Isolation of endoflagella

Flagella were obtained from B. burgdorferi by ultracentrifugation.

Immuno-electron microscopy

B. burgdorferi cells from a 100-ml culture were harvested by centrifugation at 13 000 g for 10 min and fixed in a solution of formaldehyde 0-2% w/v and glutaraldehyde 0-3% v/v in PBS for 1 h at 0°C. The samples were embedded in the low temperature resin Lowicryl K4M (Chem. Werke Lowi, Waldkraiburg, Germany). Post-embedding labelling for the immuno-electronmicroscopic localisation of p97 was performed on ultrathin sections with the protein A-colloidal gold technique. To reduce non-specific
binding, casein 1% w/v was added to the solutions containing immunoglobulins or protein A-gold. Post-staining of the sections was with uranyl acetate (pH 4.8) 4% for 3 min.

For immuno-electronmicroscopy of the isolated flagella, a grid covered with a carbon-coated Formvar film was floated on a drop of the sample. The grid was removed, dried and then incubated in serum from the immunised rabbit for 1 h. After incubation, the grid was floated on two drops of PBS containing Tween 0.05% and one drop of PBS for 2 min to remove non-specific antibodies. The grid was then floated on a drop containing a 1 in 50 dilution of 5-nm colloidal gold particles coated with protein A for 1 h. The grid was washed in two drops of PBS containing Tween 0.05% and one drop of distilled water and was finally stained by floating on a drop of phosphotungstic acid (pH 7.0) 2%. Electronmicrographs were taken with a Philips EM 301 instrument (Philips, Eindhoven, Netherlands) at calibrated magnifications.

Isolation and characterisation of the DNA of the recombinant \( \lambda \) gt11

The cultivation of the recombinant \( \lambda \) gt11 and the isolation of the bacteriophage DNA was carried out by the plate lysate method. The purified DNA was cleaved with EcoRI and the restriction fragments were analysed by agarose gel electrophoresis after staining with ethidium bromide.

Results

For the cloning of immunogenic protein sequences, DNA of \( B. burgdorferi \) was isolated and cleaved with the endonuclease EcoRI, and the DNA fragments were ligated with the EcoRI site of the bacteriophage \( \lambda \) gt11. For the screening of the resulting gene bank, the fusion proteins were expressed in \( E. coli \) strain Y1089. The hybrid proteins, expressed by the recombinant phages, consisted of \( \beta \)-galactosidase and random sequences of \( B. burgdorferi \). These were recognised in colony immuno-screening if they represented an antigenic structure of \( B. burgdorferi \). Five of c. 20 000 plaques showed a positive reaction and no reaction with the negative controls. One of these five clones was investigated further and described for this study. To obtain larger amounts of the fusion protein, the phages were subcloned in \( E. coli \) strain Y1090. The cell lysate of this clone was analysed by SDS-PAGE. An additional band of c. 200 Kda (p200) was identified after staining the \( E. coli \) lysate with Coomassie Blue, in comparison with an \( E. coli \) lysate after infection with \( \lambda \) gt11 without a \( B. burgdorferi \) insert. This band was also recognised by the human serum containing \( B. burgdorferi \) antibodies (fig. 1) but not with a negative human serum. Other bands of the lysate showed only weak reactions.

To characterise the polypeptide of \( B. burgdorferi \) present in the fusion protein, three methods were used. Firstly, the 200-Kda protein was electro-eluted from a preparative SDS-PAGE gel and added to the \( B. burgdorferi \) antiserum for absorption of the antibodies that recognised the 200-Kda protein. \( B. burgdorferi \) antigen was separated by SDS-PAGE and transferred to the PVDF membrane. The positive serum, with or without absorption by the fusion protein, was incubated with the blot. In the blot, immunostained by the pre-absorbed serum, a band of c. 97 Kda was missing, compared with the result given by the non-absorbed serum (fig. 2, lane 3). Secondly, antibodies were eluted from protein p200 bound to the blot membrane. This eluate reacted with the 97-Kda (p97) protein of \( B. burgdorferi \) in Western blots (fig. 2, lane 4). Thirdly, a rabbit was immunised with the gel-purified fusion protein. The resulting antibodies reacted in Western blots with protein p200 in the \( E. coli \) lysate, and from the blotted \( B. burgdorferi \), antigen p97 was recognised (fig. 2, lane 5). Anti-\( \beta \)-galactosidase antibodies stained only p200.

Out of 20 sera with \( B. burgdorferi \) antibodies of the IgG and IgM class, four showed strong reactions, 13
Fig. 2. Western blot of the B. burgdorferi antigen: lane 1, mol. wt marker; 2, immunoreaction of B. burgdorferi antibody-positive human serum; 3, immunoreaction of the serum used in lane 2 after absorption with protein p200; 4, immunoreaction of antibodies, eluted from protein p200; 5, immunoreaction of a rabbit serum after immunisation with protein p200 and absorption with a negative E. coli lysate.

weak reactions and three no reaction with p97 in a Western blot. The 20 negative controls did not react.

The localisation of protein p97 in the B. burgdorferi cell was studied by immuno-electronmicroscopy. Ultrathin sections of fixed and resin-embedded organisms were incubated with antibodies against p200 and p97 respectively. Bound antibodies were recognised by colloidal gold-labelled protein A. The gold particles were predominantly associated with the endoflagella in the periplasmic space (figs. 3, 4). Anti-p200 and anti-p97 antibodies showed the same staining pattern along the entire flagella. The negative controls did not show a typical distribution.

To obtain further evidence for a flagellar localisation of the cloned antigen, flagella were isolated by the method of Coleman and Benach. Firstly the flagella were investigated by immuno-electronmicroscopy, by means of immune rabbit serum or with the isolated anti-p97 antibodies (fig. 5). The serum of the rabbit before immunisation served as a negative control. The antibodies, in contrast to the negative control, became bound along the entire flagellum. Secondly, after staining with Coomassie Blue, the Western blot of the purified flagella revealed two distinct bands, one intense, representing the major flagellar protein p41, and a second of less intensity, with M, of c. 97 Kda. The serum of the immunised rabbit, as well as the eluted human anti-p97 antibodies, stained the 97-Kda protein only (fig. 6).

For its characterisation, the DNA of the recombi-

Fig. 3. Ultrathin sections of B. burgdorferi cells with p200 antibodies and protein-A gold complexes. Bar = 100 nm. (A) Negative control (F, flagella; CM, cytoplasmic membrane; OM, outer membrane). (B) Labelling with rabbit-IgG directed against p200. Arrows indicate immunogold particles in the periplasmic space associated with the flagellar bundle.

nant bacteriophages was isolated and cleaved with the endonuclease EcoRI. The resulting fragments were analysed by agarose gel electrophoresis. DNA fragments of the two $\lambda$gt11 arms of 19-6 kb and 24-1 kb and an additional DNA of about 6-6 kb in length could be recognised (fig. 7).

Fig. 4. Ultrathin sections of B. burgdorferi cells labelled with p97 antibodies and protein-A gold-complexes. Bar = 450 nm. (A) Negative control with a B. burgdorferi antibody-negative human serum. (B) Labelling with p97 antibodies eluted from a blot membrane. Arrows indicate typical attachment sites of the gold marker, associated with the flagella.
Discussion

The motility of the helical \( B. \) burgdorferi is brought about by c. 10 flagella, located in the periplasmic space between the cytoplasmic membrane and the outer envelope.\(^{27}\) Recently, the major flagellar protein p41 was characterised and the nucleotide sequence, coding for 336 amino acids, was determined. The sequence data demonstrated a high homology to the 33-Kda flagellar protein of \( T. \) pallidum.\(^8\) Coleman and Benach isolated \( B. \) burgdorferi flagella by CsCl gradient centrifugation.\(^9\) In the SDS-PAGE of the product, stained with Coomassie Blue, protein p41 and a minor band of c. 85 Kda were identified. The authors suggested that this 85-Kda protein might be a dimer of protein p41, but it is also possible that the protein was a different structural part of the flagella. It remained unclear whether the flagella in \( B. \) burgdorferi consisted of only p41 subunits, or whether it was assembled from several structural proteins. Flagella of other spirochaetes, e.g., some treponemes or leptospires, consist of different proteins.\(^{28-31}\)

In the present study, an antigenic hybrid protein was expressed in \( E. \) coli, by means of DNA fragments of \( B. \) burgdorferi ligated with the DNA of a \( \lambda \) gt11 bacteriophage. The resulting protein had an \( M_\text{r} \) in the range of 200 Kda. The mass of \( \beta \)-galactosidase in the fusion protein was 114 Kda, so that the expressed borrelia polypeptide had an \( M_\text{r} \) of c. 90 Kda. This was the minimum mass that the corresponding protein from \( B. \) burgdorferi could have. The native protein was identified by several methods as having an \( M_\text{r} \) of 97 Kda (p97). Antibodies directed against the hybrid protein p200 or the corresponding protein p97 of \( B. \) burgdorferi, characterised and identified in this report, reacted with antigens located in the region of the periplasmic flagella as shown by immuno-electron-microscopy of the bacterial cells. The immunogold complexes were distributed along the total length of the flagella (as shown by longitudinal sections of borrelia cells marked with immunogold complexes and "on grid" immunolabelling of the isolated flagella) and at the position of the flagellar bundle in the periplasmic space (as shown by transverse sections). The results obtained from immuno-electronmicro-
scopy of purified flagella also indicated that protein p97 was associated with the flagella. This evidence was impressively supported by the Western blot of the flagellar fraction.

The experimental data of the blots further suggested an absence of antigen structures common to protein p97 and the major flagellar protein p41. The latter was not a constituent of p97, and p97 might, therefore, have been a structural entity of the whole flagella, but to a lesser extent than p41, as indicated by results of SDS-PAGE. It might be speculated that protein p97 is necessary for preservation of the flagellar structure. Similar proposals were made for the two flagellar proteins found in *Treponema phagedenis* and other types of spirochaetal flagella which consist of different proteins.18

In sera of patients with clinically and serologically diagnosed Lyme disease, p97 antibodies, unlike p41 antibodies, could not always be detected. The p97 antibodies seemed to be specific, because the negative controls showed no reaction in the Western blots. It is still unclear whether this protein p97 was expressed by different *B. burgdorferi* strains. Various *B. burgdorferi* strains show proteins in the range of 90-100 KDa with different concentrations.8,9,32 In the *B. burgdorferi* strain we used, protein p97 was not a dominant protein like the surface proteins Osp A, Osp B, the common antigen or the flagellar protein p41.

Further studies are needed to demonstrate whether protein p97 differs in different *B. burgdorferi* strains and whether it is useful in diagnostic tests. More information about p97 would become available through a better characterisation of its gene. The determination of the nucleotide sequence is under investigation.

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References