Detection of *Borrelia burgdorferi* in cerebrospinal fluid by the polymerase chain reaction

W. H. KRÜGER* and M. PULZ

Institute of Medical Microbiology and Immunology, University Hospital Eppendorf, Martinistrasse 52, 2000 Hamburg 20, Germany

Summary. The polymerase chain reaction (PCR) was used to amplify specific DNA sequences from different clinical isolates of *Borrelia burgdorferi* and from cerebrospinal fluid (CSF) of two patients with Lyme disease of the central nervous system. The amplification products were separated by polyacrylamide gel electrophoresis and visualised by ethidium bromide staining. The definitive identification of amplified DNA as a part of the *B. burgdorferi* flagellin gene was achieved by hybridisation to a 40-base oligonucleotide probe complementary to a part of the spirochaetal gene but not to the primers. Attempts to cultivate borreliae from either patient were unsuccessful and one patient had no serological marker in serum or CSF to indicate borreliosis. Clinical symptoms of both patients regressed with antibiotic therapy. The PCR system is a powerful and rapid technique to amplify flagellin gene sequences from CSF of patients with neuroborreliosis. Only one-tenth of the time needed for cultivation was required from CSF sampling to diagnosis. Gene amplification might, for the first time, allow effective monitoring of therapy for patients with Lyme disease of the central nervous system.

Introduction

Lyme borreliosis, caused by the spirochaete *Borrelia burgdorferi* is the most frequent tick-borne disease in Europe and the USA.1-4 Meningitis, cranial neuritis and painful radiculoneuritis have been reported to be common neurological disorders in this infection.5-8 Diagnosis of borreliosis of the central nervous system relies mainly on clinical and serological features. Antibodies against *B. burgdorferi* in blood and cerebrospinal fluid (CSF) are demonstrable by indirect immunofluorescence, ELISA and Western blotting.9-11 Complete confidence in these tests is probably misplaced; around 20–50% of cases have no measurable antibody response during the first few weeks12-15 and cases of seronegative neuroborreliosis have been described in which *B. burgdorferi* was isolated from CSF.4,15-17 However, cultivation of *B. burgdorferi* from CSF requires 4–6 weeks and is often unsuccessful.18-20 Thus, techniques for the rapid detection of spirochaetes in clinical samples are needed.21,22

The polymerase chain reaction (PCR) is a technique for detecting nucleic acids based on a nearly exponential multiplication of a defined DNA template.23 In this report, we describe two oligonucleotides designed to amplify a segment of the *B. burgdorferi* flagellin gene. A further oligonucleotide, with no homology to the PCR primers, was used for the specific detection of the amplified products by hybridisation. We used this method to detect Lyme disease spirochaete DNA in CSF from two patients with neuroborreliosis.

Materials and methods

**Bacterial strains**

*B. burgdorferi* reference strains B31 (ATCC 35210) and IRS (ATCC 35211) were used as positive controls. Clinical isolates of *B. burgdorferi* from CSF (strains 50 and 387) and from tissue biopsy samples of erythema chronicum migrans (strains N34 and PKO) were kindly provided by R. Ackermann, Cologne (N34), V. Sticht-Groh, Würzburg (50, 387) and B. Wilske, Munich (PKO). All borreliae were cultured in modified Kelly medium and passaged twice weekly.24 Numbers of spirochaetes were determined by dark field microscopy with a Petroff-Hauser counting chamber.

**Patients**

The first patient was a 50-year-old man. In November 1989 he had a febrile illness. In December he developed neurological symptoms including meningitis and cranial nerve (VI and VII) palsies. He was admitted to our hospital with the additional signs of an early psychosyndrome. He remembered no tick...
bite. The second patient was a 10-year-old boy who was admitted to hospital because of a left sided facial palsy. The parents reported a tick bite followed by an apparent erythema migrans sometime before the onset of the neurological disorder.

**Laboratory investigations and clinical specimens**

Examination of CSF from the first patient showed a lymphocytosis (4200 cells/μl) and elevated total protein. IgG and glucose concentrations. Oligoclonal IgG was detected in CSF by iso-electric focusing. The CSF albumin:serum albumin ratio revealed a blood-brain barrier dysfunction. *B. burgdorferi*-specific Western blots showed a negative result with serum but revealed a specific IgG response in CSF. Cultivation of *B. burgdorferi* from CSF samples was attempted in duplicate before antibiotic therapy was started but cultures remained negative after incubation for 6 weeks. During therapy with cefotaxime (2 g, three times daily for 14 days) cranial neuritis disappeared and lymphocytosis decreased.

Examination of CSF from the second patient also revealed a lymphocytosis. *Borrelia*-specific antibodies were not detected in serum or CSF and spirochaetes were not cultivated from CSF. The diagnosis was based primarily on clinical features. Lymphocytosis and paresis regressed with antibiotic therapy (penicillin-G, 4-5 million units, three times daily for 14 days) cranial neuritis disappeared and lymphocytosis decreased.

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Neurosyphilis involving the CNS was excluded in both patients by serological tests. Meningitis, caused by classical bacterial agents, and tuberculosis were excluded by standard examinations. Cell culture and serological screening for neurotropic viruses gave negative results.

CSF samples were used for the PCR.

**Negative controls**

As negative controls for PCR, DNA extracted from the human lymphocyte line H9-(a gift from R. Gallo, Bethesda, MD, USA) and CSF samples from neurosurgical patients with no evidence of Lyme borreliosis were used. PCR assays were also performed with suspensions (10^5 bacteria/ml) of agents of typical bacterial meningitis—*Escherichia coli*, *Neisseria meningitidis*, *Haemophilus influenzae*, *Streptococcus pneumoniae*, *Staphylococcus aureus* and *Mycobacterium tuberculosis*.

**Synthetic oligonucleotides**

The oligonucleotides were synthesised on an Applied Biosystems synthesiser with the automated phosphoramidite coupling method. The primer pair for the PCR procedure, termed WK1 and WK2, and the hybridisation oligonucleotide WK3 were deduced from the gene sequence for the flagellin of *B. burgdorferi* published by Gassmann et al.25 Nucleotide sequences are shown in fig. 1. The oligonucleotide used in the hybridisation assay was radiolabelled with [γ^32P]ATP.26

**PCR**

Cultures of *B. burgdorferi* containing c. 10^8 organisms/ml were diluted up to 10^5-fold in distilled water. The concentration of H9—DNA was adjusted to 0.1 μg/μl. CSF samples, c. 3 ml, were centrifuged at 1000 g for 10 min, then the pellet was resuspended in a few drops of supernate. All samples, except for the H9—DNA, were boiled for 10 min in a water bath. Amplifications were performed with 10 μl of each sample added to a final volume of 100 μl with 10 μl of 10-fold concentrated *Thermus aquaticus* polymerase reaction buffer, dNTP mix, distilled water and 2U of *T. aquaticus* polymerase (Perkin-Elmer-Cetus) as described previously.26 After a primary denaturing process for 5 min at 95°C to ensure complete denaturation of target DNA, the samples were subjected 35 times in succession to the following thermal cycle: denaturation for 1 min at 95°C, annealing for 1 min at 48°C and polymerisation for 2 min at 72°C. Thermal cycling was performed in a programmable heat block (Perkin-Elmer-Cetus).26

**Analysis of amplification products**

Amplified samples (50-μl volumes) were subjected to electrophoresis through polyacrylamide (7-5%) gel and visualised by staining with ethidium bromide. The nucleic acids were transferred from the polyacrylamide gel to Zeta Probe membranes (BioRad) by electroblotting. Prehybridisation was performed in 5 x SSPE (0.9 M NaCl, 50 mM NaH2PO4 and 5 mM EDTA, pH 7-4), 5 x DET (polyvinylpyrrolidone, Ficoll, bovine serum albumin, each at 0.1% w/v) and sodium dodecyl sulphate 0.2% w/v for 2 h at 55°C. Radiolabelled oligonucleotides, 10^5 cpm/ml, were then added and hybridisation was allowed to proceed for 20 h at 55°C in the same buffer. The membranes were then washed twice in 5 x SSPE-sodium dodecyl sulphate 0.2% w/v at 30°C for 10 min each time and once for 10 min at 55°C in the same solution. The membranes were dried in air and autoradiographed at −80°C.26,27

**Results**

**Sensitivity and specificity of primers for detection of *B. burgdorferi* flagellin gene**

The primers were tested for PCR amplification of nucleic acids of *B. burgdorferi*, human lymphocytes

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**Fig. 1. Oligonucleotides used for amplification and hybridisation with *B. burgdorferi* flagellin gene sequences.**

<table>
<thead>
<tr>
<th>Primer</th>
<th>Sequence</th>
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<tbody>
<tr>
<td>WK1</td>
<td>AAGGAATGGGCAGTTCAATC (271-290)</td>
</tr>
<tr>
<td>WK2</td>
<td>GTTCTACTTCGATAACGACT (541-560)</td>
</tr>
<tr>
<td>WK3</td>
<td>AATGCACATGGTTACAAAACATCTGTC (396-425)</td>
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and typical bacterial agents of meningitis. The amplified 290-bp fragment was visible in ethidium bromide-stained gels when DNA of any *B. burgdorferi* isolate was the target and the spirochaete suspension was diluted up to 10⁵-fold, equivalent to c. 10 bacteria per sample (fig. 2). This calculation was based on a bacterial density of 10⁸ spirochaetes/ml of culture fluid, determined with a Petroff-Hauser counting chamber. A visible amplification product of c. 290 bp hybridised with the oligonucleotide probe WK3. We could not amplify any DNA sequences from human lymphocytes. The hybridisation with probe WK3 was specific to DNA fragments of the expected size in positive samples. There was no evidence for non-specific hybridisation of probe WK3 to human DNA. Nucleic acid sequences of bacteria other than *B. burgdorferi* were not amplified with the described oligonucleotides.

Detection of *B. burgdorferi* DNA in clinical samples

After PCR amplification, electrophoresis and hybridisation, the presence of the *B. burgdorferi* flagellin gene was demonstrated in culture samples of different *B. burgdorferi* strains and in CSF samples from both patients with clinically diagnosed Lyme borreliosis (fig. 3). All control CSF samples gave negative results both with ethidium bromide staining and with hybridisation.

Discussion

The cultivation of *B. burgdorferi* has been considered to be the reference method for demonstrating the infectious agent of Lyme disease in clinical specimens from patients and in ticks. However, culture of this fastidious organism is difficult and requires incubation for about 6 weeks.¹⁷,¹⁹,²⁰ We have established PCR for rapid and sensitive detection of *B. burgdorferi* in CSF. Because clinical samples contain only a few organisms, sensitivity is an obligate requirement for any PCR assay for the Lyme disease spirochaete. This system is practicable because we were able to detect a few (≤10) bacteria in dilutions of liquid cultures. DNA was amplified from all *B. burgdorferi* isolates tested. By employing larger sample volumes, the sensitivity of PCR for clinical use might be increased.

The gene amplification technique offers advantages over culture. It is not necessary to have viable organisms in specimens. Thus, in our second patient with neuroborreliosis, the clinical diagnosis of Lyme disease could be confirmed only by gene amplification. Cultures remained negative and borrelia-specific antibodies were not detected. Cases with positive CSF cultures for *B. burgdorferi* but with negative serology in CSF and serum have been described. In the past, the diagnosis in such cases depended upon successful culture or subsequent seroconversion and was very time-consuming. PCR allows a rapid diagnosis, even in seronegative patients.¹⁰,¹⁶–¹⁸,²⁰

Survival of *B. burgdorferi* in antibiotic-treated patients with Lyme disease of the central nervous system has been described.¹⁷ We were able to detect borrelia-specific DNA in CSF from the first patient 2 weeks after beginning antimicrobial therapy; 4 weeks later, *B. burgdorferi* DNA was not detectable in CSF samples, in parallel with the clinical improvement seen. Thus PCR may prove to be a useful and sensitive method for monitoring therapy in patients with neuroborreliosis.

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Fig. 2. Electrophoretic separation (a) and corresponding Southern blot hybridisation assay (b) of amplified *B. burgdorferi* flagellin gene sequences. Lanes 1 and 2: negative controls, no signal was detected when distilled water (1) or human lymphocyte DNA (2) was submitted to PCR; 3–7: amplified DNA from 10-μl samples of *B. burgdorferi* strain B31 cultures, diluted 10³–10⁵-fold respectively.
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Fig. 3. Amplified flagellin gene sequences from culture dilutions (10⁻²) of different B. burgdorferi strains and from CSF. The electrophoretic profiles (a) and corresponding Southern blot patterns (b) are shown after hybridisation with the oligonucleotide probe WK3. Control preparation (left lane) for size in bp was pBR322 DNA, digested with HaeIII. Lane 1: distilled water; 2: human DNA from H9- cells (negative control); 3: B. burgdorferi strain B31 (ATCC 35210); 4: strain IRS (ATCC 35211); 5-8: clinical isolates 50, 387, N34 and PKO respectively; 9: CSF sample from patient 1; 10: CSF sample from patient 2.

This test may also help detect B. burgdorferi in other specimens, e.g., synovial fluid from patients with Lyme arthritis, a manifestation of B. burgdorferi infection more frequently reported from the USA. The cultivation of B. burgdorferi from synovial fluid appears to be even less successful than from CSF.

We thank Carola Schlüter for preparing the photographs.

References


