Real time polymerase chain reaction: a new powerful tool for the diagnosis of neurobrucellosis

J D Colmenero, M I Queipo-Ortuño, J M Reguera, G Baëza, J A Salazar, P Morata


**Background/methods:** We compared the diagnostic yield of a real time polymerase chain reaction (PCR) assay in cerebrospinal fluid (CSF) samples with conventional microbiological techniques for the diagnosis of neurobrucellosis. Following amplification of a 223 bp sequence specific for *Brucella* genus, melting curve analysis was performed to verify the specificity of the PCR products.

**Results:** All six patients with neurobrucellosis (three meningitis and three meningoencephalitis) had a positive real time PCR assay, whereas CSF cultures and Wright seroagglutination tests were positive in only two and four cases, respectively. *Brucella* specific amplicons were easily demonstrated by their characteristic melting temperature in all the real time PCR assays.

**Conclusion:** LightCycler based real time PCR assay in CSF samples is more rapid and sensitive than conventional microbiological tests. This technique could be useful for the rapid diagnosis of neurobrucellosis.

**patients and methods**

**Patients**

From January 1998 to December 2002, six patients with neurobrucellosis were diagnosed, treated, and followed at the Infectious Diseases Unit or Neurology Service of Carlos Haya University Hospital, Spain. The diagnosis of brucellosis was established according to one of the following criteria: first, isolation of *Brucella* spp in blood or any other body fluid or tissue sample, or, second, the presence of a compatible clinical picture together with the demonstration of specific antibodies at significant titres or seroconversion. Significant titres were considered to be a Wright’s seroagglutination ≥1/160 or a Coombs antibrucella test ≥1/320. Neurobrucellosis was diagnosed in patients with neurological symptoms and signs not otherwise explainable and at least one of the following criteria: (a) isolation of *Brucella* from CSF; (b) demonstration of antibodies to *Brucella*, agglutinating or non-agglutinating, in CSF (at any titre) in the presence of one or more abnormalities of the CSF (>10 cells/μl, protein levels >45 mg/dl, or glucose levels <40 mg/dl or 40% of the concomitant blood glucose level).

**Specimen collection and processing**

Samples of CSF were cultured onto blood and chocolate agar media, MacConkey agar, and *Brucella* agar (Biomedics, San Sebastian de los Reyes, Madrid, Spain). In order to rule out the possibility of mycobacterial and fungal infection all samples were also cultured onto Lowestein and Saburaud medium. The plates were incubated in a 10% CO₂ atmosphere at 37°C for at least 7 days. If growth appeared, the suspected colonies were identified by colonial morphology, Gram staining, oxidase, catalase and urease tests, and positive agglutination with specific antiserum.

**DNA extraction**

CSF samples were maintained at −20°C until processing. DNA was extracted, by boiling, from 200 μl to 1 ml of CSF placed in a 1.5 ml microcentrifuge tube and centrifuged for 15 min at 15 000 X g. The supernatant was eliminated and the pellet resuspended in 40 μl of sterile water and subjected to boiling in a water bath for 10 min, again eliminated and the pellet resuspended in 40 μl of sterile water and subjected to boiling in a water bath for 10 min, cooled on ice and centrifuged for 10 s at 15 000 X g. The supernatant was again eliminated and the pellet resuspended in 40 μl of sterile water and subjected to boiling in a water bath for 10 min, cooled on ice and centrifuged for 10 s at 15 000 X g, before storing at −20°C until use. A total of 2 μl of the suspension was used for PCR.

**Real time PCR with SYBR Green I**

Primers were selected from the conserved region of the gene that encodes an immunogenic membrane protein of 31 kDa of *Brucella abortus* specific to the *Brucella* genus. The pair of 21 nucleotide primers B₄ (5’ TGG CTC GGT TGC CAA TAT CAA 3’) and B₅ (5’ CGC GCT TGC CTT TCA GGT CTG 3’) (Tib Molbiol, Berlin, Germany) were used in the amplification process. PCR amplifications were performed in capillary tubes with 20 μl of final volume in a LightCycler instrument (Roche Diagnostic, Mannhein, Germany). Reaction mixtures contained 2 μl of LightCycler FastStart DNA master mixture for SYBR Green I (Roche Diagnostic, Germany), 0.5 μM each primer, 4 mM MgCl₂, and 2 μl of DNA template. All capillaries were sealed and then centrifuged at 500 X g for 1.5 min at 50°C. The mixture was then cooled on ice and stored at −20°C until use. To validate the sensitivity, 100-fold dilutions of PCR products were used as templates.

**Abbreviations:** CNS, central nervous system; CSF, cerebrospinal fluid; CT, computed tomography; MR, magnetic resonance; PCR, polymerase chain reaction

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*Brucellosis* is a worldwide zoonosis. In many countries, especially around the Mediterranean basin, and in the Middle East, India, Mexico, and Central and South America, the disease remains endemic.1

Human brucellosis is a systemic infection with a wide clinical spectrum. From 20–40% of patients with brucellosis present focal complications, which can affect any organ or system,2 including the central nervous system (CNS), where it shows great clinical polymorphism. Although the incidence of CNS involvement is not high (1–2% of all cases) it results in considerable morbidity and mortality.2–4 Neurobrucellosis is often difficult to diagnose because the titres of specific antibrucella antibodies in cerebrospinal fluid (CSF) are usually low and the yield of the cultures is poor, with rates of positivity of just 40–50%.5

Among molecular techniques, the polymerase chain reaction (PCR) has already proved to be a very useful tool for the diagnosis of many CNS infections and is now considered the gold standard in some.6,7 We describe a SYBR Green I LightCycler based real time PCR assay in CSF that encodes an immunogenic membrane protein of 31 kDa specific to the *Brucella* genus. The pair of 21 nucleotide primers B₄ (5’ TGG CTC GGT TGC CAA TAT CAA 3’) and B₅ (5’ CGC GCT TGC CTT TCA GGT CTG 3’) (Tib Molbiol, Berlin, Germany) were used in the amplification process. PCR amplifications were performed in capillary tubes with 20 μl of final volume in a LightCycler instrument (Roche Diagnostic, Mannhein, Germany). Reaction mixtures contained 2 μl of LightCycler FastStart DNA master mixture for SYBR Green I (Roche Diagnostic, Germany), 0.5 μM each primer, 4 mM MgCl₂, and 2 μl of DNA template. All capillaries were sealed and then centrifuged at 500 X g for 1.5 min at 50°C. The mixture was then cooled on ice and stored at −20°C until use. To validate the sensitivity, 100-fold dilutions of PCR products were used as templates.
Stranded PCR product was measured during the 72 °C extension step by detection of fluorescence associated with the binding of SYBR green to the product. For data analysis, baseline adjustment was carried out in “none” mode and fluorescence curve analyses were carried out in the “fit points” mode of the LightCycler software. Positive controls based on DNA from B abortus B-19 were included in all the tests, as were negative controls containing all the elements of the reaction mixture except template DNA. To guarantee the reliability of the results all the samples were processed in duplicate. To prevent contamination, universal precautions were exercised and one-way flow of DNA extraction and amplification were used. To avoid potential subjectivity, the status of each patient for Brucella infection was unknown during the PCR assay. The test was only considered positive if the signal from the amplified fragment was clearly visible in both samples.

RESULTS

Five of the patients were men and one a woman. Their mean age was 37.6 ± 16.6 years (range: 21–69 years). Four patients had usual contact with sheep or goats, one habitually consumed unpasteurised dairy products, and the other had no known prior exposure to Brucella infection.

Three patients had meningoencephalitis and three had meningitis. The duration of the symptoms prior to diagnosis was 90 ± 138 days (range: 5–365 days). Four patients (66.6%) presented fever and only three (50%) had clear signs of meningeval irritation. Other relevant clinical data are shown in Table 1.

Examination of CSF showed findings compatible with meningitis in all cases (Table 2). Antibruccella antibodies could also be demonstrated in CSF in all cases. Wright seroagglutination was positive in four cases (66.6%) and in the other two cases non-agglutinating antibodies were shown by Coombs antibruccella test. In fact, Coombs antibruccella titres were higher than those of the agglutination test (Table 2).

CSF IgG values were increased in all cases (range: 2.34–6.42 g/L), and CSF protein electrophoresis showed oligoclonal bands in the three cases in who it was carried out.

Five patients had brain magnetic resonance (MR) imaging studies and the remaining patient a cranial computed tomography (CT) scan. In cases 1, 2, and 6, one or more hyperintense lesions on T2 weighted images were seen in periventricular white matter, the right cerebellum, and the right occipital hemisphere, respectively. In the remaining three cases, neuroimaging studies were normal. Brucella was isolated from CSF in two cases, whereas real time PCR assay was positive in all six cases and in each case Brucella specific amplicons could be easily distinguished by their characteristic melting temperature.

DISCUSSION

The clinical spectrum of neurobrucellosis is very heterogeneous. Different clinical pictures have been described, including meningitis, meningoencephalitis, intracerebral haemorrhage, benign intracranial hypertension, optic neuritis, arachnoiditis, polyradiculoneuritis, myelitis, and various combinations of these. Accordingly, neurobrucellosis, like neurosyphilis, is easily confused with many other neurological, neurosurgical, or even psychiatric processes.

As seen in this and other studies, although the meninges are almost always involved in patients with neurobrucellosis, the clinical signs of meningeal irritation are often lacking or, if present, may well be discrete. Furthermore, the fact that CSF cultures are negative in over 50% of cases, that IgG oligoclonal bands are often present, and that these patients are not always seen by specialists in infectious diseases, explains why neurobrucellosis is not always considered in the differential diagnosis.

Adequate antibiotic treatment usually results in a good prognosis for patients with neurobrucellosis. However, 20–30% of patients still present neurological sequelae, which are clearly related with the delay in diagnosis and initiation of correct antibiotic treatment.

Our group has reported that PCR based methods provide better results than conventional microbiological techniques for the diagnosis of both primary infection and relapses of brucellosis. Real time PCR is a new automated amplification technique that quantitatively monitors PCR products as they accumulate during thermal cycling. LightCycler technology (Roche Diagnostic, Germany) combines rapid amplification and sequence-specific detection of amplicons in an automated and standardized format. Because this probe does not require post-amplification handling, much faster assays are possible.

This study showed real time PCR assay to be positive in all six cases of neurobrucellosis, whereas the sensitivity of

<table>
<thead>
<tr>
<th>Table 1 Summary of clinical finding</th>
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</thead>
<tbody>
<tr>
<td><strong>Patient 1</strong></td>
</tr>
<tr>
<td>----------------</td>
</tr>
<tr>
<td><strong>Duration of the symptoms</strong></td>
</tr>
<tr>
<td><strong>Fever</strong></td>
</tr>
<tr>
<td><strong>Vomiting</strong></td>
</tr>
<tr>
<td><strong>Headache</strong></td>
</tr>
<tr>
<td><strong>Stiff neck</strong></td>
</tr>
<tr>
<td><strong>Obnubilation</strong></td>
</tr>
<tr>
<td><strong>Papilloedema</strong></td>
</tr>
<tr>
<td><strong>Impaired vision</strong></td>
</tr>
<tr>
<td><strong>Hemiparesis</strong></td>
</tr>
<tr>
<td><strong>Deafness</strong></td>
</tr>
<tr>
<td><strong>Seizures</strong></td>
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<tr>
<td><strong>Ataxia</strong></td>
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</tbody>
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seroagglutination and cultures of CSF samples was 66.6% and 33.3%, respectively. Real time PCR was negative in all the control samples, which is in keeping with the specificity of the primers used. Previous studies have demonstrated the high specificity of the B4 and B5 primers with a wide panel of microorganisms.13

The LightCycler assay used is rapid, easy, and objective, allowing sensitive and specific identification of PCR products in less than 2 hours. Moreover, the technique is a closed-tube system; this considerably reduces the risk of contamination by product carryover and allows simultaneous handling of a large number of samples, making it very suitable for use in a clinical laboratory. Finally, it should not be forgotten that Brucella spp are class III pathogens requiring special protection measures. PCR assays completely obviate the need for direct handling of the pathogen, thus greatly reducing the risk of infection of laboratory personnel.

In conclusion, although the results of this study are preliminary, the high sensitivity and specificity of this real time PCR assay, together with its speed and versatility, make it a very useful tool and could lead to it being considered the new gold standard for the diagnosis of neurobrucellosis.

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Competing interests: none declared

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REFERENCES

Table 2 Biological and microbiological finding in CSF samples of patients with neurobrucellosis

<table>
<thead>
<tr>
<th>Glucose mmol/L</th>
<th>Patient 1</th>
<th>Patient 2</th>
<th>Patient 3</th>
<th>Patient 4</th>
<th>Patient 5</th>
<th>Patient 6</th>
</tr>
</thead>
<tbody>
<tr>
<td>CSF/plasma glucose ratio (%)</td>
<td>4.49</td>
<td>2.55</td>
<td>1.05</td>
<td>0.61</td>
<td>1.66</td>
<td>1.27</td>
</tr>
<tr>
<td>CSF WBC ml (% lymphocytes)</td>
<td>8 (75%)</td>
<td>85 (90%)</td>
<td>825 (70%)</td>
<td>579 (75%)</td>
<td>540 (70%)</td>
<td>45 (95%)</td>
</tr>
<tr>
<td>Protein g/L</td>
<td>0.85</td>
<td>3.03</td>
<td>5.35</td>
<td>0.84</td>
<td>0.80</td>
<td>1.80</td>
</tr>
<tr>
<td>Real time PCR</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Adenosine deaminase U/L</td>
<td>ND</td>
<td>11</td>
<td>35</td>
<td>17</td>
<td>7</td>
<td>14</td>
</tr>
<tr>
<td>Culture</td>
<td>–</td>
<td>B melitensis</td>
<td>–</td>
<td>–</td>
<td>B melitensis</td>
<td>–</td>
</tr>
<tr>
<td>Seroagglutination*</td>
<td>20</td>
<td>320</td>
<td>–</td>
<td>–</td>
<td>40</td>
<td>80</td>
</tr>
<tr>
<td>Coombs test*</td>
<td>160</td>
<td>5120</td>
<td>1280</td>
<td>640</td>
<td>160</td>
<td>5120</td>
</tr>
</tbody>
</table>

*Expressed as reciprocal of titre.

CSF, cerebrospinal fluid; ND, not done; PCR, polymerase chain reaction; WBC, white blood count.
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