The ligase chain reaction as a primary screening tool for the detection of culture positive tuberculosis

T M O’Connor, S Sheehan, B Cryan, N Brennan, C P Bredin

Abstract

Background—The ligase chain reaction assay uses ligase chain reaction technology to detect tuberculosis DNA sequences in clinical specimens. A study was undertaken to determine its sensitivity and specificity as a primary screening tool for the detection of culture positive tuberculosis.

Methods—The study was conducted on 2420 clinical specimens (sputum, bronchoalveolar lavage fluid, pleural fluid, urine) submitted for primary screening for *Mycobacterium tuberculosis* to a regional medical microbiology laboratory. Specimens were tested in parallel with smear, ligase chain reaction, and culture.

Results—Thirty nine patients had specimens testing positive by the ligase chain reaction assay. Thirty two patients had newly diagnosed tuberculosis, one had a tuberculosis relapse, three had tuberculosis (on antituberculous therapy when tested), and three had healed tuberculosis. In the newly diagnosed group specimens were smear positive in 21 cases (66%), ligase chain reaction positive in 30 cases (94%), and culture positive in 32 cases (100%). Using a positive culture to diagnose active tuberculosis, the ligase chain reaction assay had a sensitivity of 93.9%, a specificity of 99.8%, a positive predictive value of 93.8%, and a negative predictive value of 99.9%.

Conclusions—This study is the largest clinical trial to date to report the efficacy of the ligase chain reaction as a primary screening tool to detect *Mycobacterium tuberculosis* infection. The authors conclude that ligase chain reaction is a useful primary screening test for tuberculosis, offering speed and discrimination in the early stages of diagnosis and complementing traditional smear and culture techniques.

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Keywords: ligase chain reaction; smear; culture; tuberculosis

Tuberculosis in Ireland causes significant morbidity and mortality with a notification rate of 15/100 000 of population reported for 1994.1 One aspect of tuberculosis control that clearly needs improvement is the speed with which the diagnosis is confirmed. If individuals with pulmonary tuberculosis can be identified rapidly, tuberculosis control could potentially be improved by earlier institution of drug treatment, isolation procedures, and contact investigation.

The role of nucleic acid amplification technology in the routine diagnosis of infectious disease is rapidly being established. The use of polymerase chain reaction (PCR) techniques for the detection of *Mycobacterium tuberculosis* has a reported sensitivity of 55–100%.2 However, contamination of specimens with PCR products can lead to false positive results, and inhibitors of PCR can lead to false negative results.3 The ligase chain reaction is an in vitro nucleic acid amplification technique that exponentially amplifies selected DNA sequences.4 The Abbott ligase chain reaction assay was first introduced in 1995 for the diagnosis of *Chlamydia trachomatis* genital-urinary infection.4 More recently it has been used for the direct detection of *M tuberculosis* in respiratory specimens. Its specificity and sensitivity have been reported as 74–90% and 100%, respectively.5,6 Chemical inactivation of amplification product minimises the risk of nucleic acid contamination of the specimen and consequent false positive results. However, the ligase chain reaction assay may be oversensitive, as evidenced by positive results in patients with previously treated tuberculosis that is now healed. We have previously reported in a small series its suitability as a primary screening method for testing all samples for the presence of *M tuberculosis*,8 and have since performed assays on all samples sent for *M tuberculosis* smear and culture testing to a regional medical microbiology laboratory. We undertook this study to evaluate its sensitivity and specificity in culture positive tuberculosis.

Methods

STUDY SUBJECTS

During the period from 1 April 1997 to 1 June 1998 a total of 2420 clinical specimens (sputum, bronchoalveolar lavage fluid, pleural fluid, urine) were submitted for tuberculosis investigations to the regional medical microbiology laboratory, Cork University Hospital. Specimens were tested in parallel by smear, ligase chain reaction, and culture. The catchment area was the south-west region of the Republic of Ireland (Counties Cork and Kerry) with a population of 546 000.

STUDY DESIGN

The microscopy procedure consisted of initial auramine stain for screening, with positive samples being confirmed by Ziehl-Neelsen
Table 1  Positive tests according to clinical diagnosis (n=39) from April 1997 to June 1998

<table>
<thead>
<tr>
<th>Diagnosis</th>
<th>Smear positive</th>
<th>Culture positive</th>
<th>Ligase chain reaction positive</th>
</tr>
</thead>
<tbody>
<tr>
<td>New TB (n=32)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sputum (n=20)</td>
<td>21 (66%)</td>
<td>32 (100%)</td>
<td>30 (94%)</td>
</tr>
<tr>
<td>BAL fluid (n=7)</td>
<td>17 (85%)</td>
<td>20 (100%)</td>
<td>19 (95%)</td>
</tr>
<tr>
<td>Pleural fluid (n=3)</td>
<td>3 (43%)</td>
<td>7 (100%)</td>
<td>7 (100%)</td>
</tr>
<tr>
<td>Urine (n=2)</td>
<td>0 (0%)</td>
<td>3 (100%)</td>
<td>2 (67%)</td>
</tr>
<tr>
<td>Relapse TB</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sputum (n=1)</td>
<td>1 (100%)</td>
<td>1 (100%)</td>
<td>1 (100%)</td>
</tr>
<tr>
<td>TB on treatment</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sputum (n=3)</td>
<td>0 (0%)</td>
<td>0 (0%)</td>
<td>3 (100%)</td>
</tr>
<tr>
<td>Healed TB</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sputum (n=3)</td>
<td>0 (0%)</td>
<td>0 (0%)</td>
<td>3 (100%)</td>
</tr>
</tbody>
</table>

The distribution of specimen type in the group with newly diagnosed tuberculosis is shown. All patients in the other groups had pulmonary tuberculosis. BAL = bronchoalveolar lavage.

The clinical records of all patients with specimens which were positive on smear, ligase chain reaction, or culture assays for M tuberculosis during a 14 month period (April 1997 to June 1998) were reviewed and their clinical details recorded. Patients were described as having newly diagnosed tuberculosis, relapse of tuberculosis, recently diagnosed tuberculosis (on antituberculosis treatment at the time of analysis), or old (healed) tuberculosis as determined by their clinicians. Therapeutic decisions were not made on the basis of ligase chain reaction results. In the case of multiple specimens taken from one patient simultaneously, the various tests were considered positive if at least one of the specimens was positive. However, there were no discrepancies noted between multiple specimens in terms of their smear, ligase chain reaction, or culture results.

**STATISTICAL ANALYSIS**

The results were calculated using the Graphpad Instat statistical software package (San Diego, CA, USA) to determine the sensitivity, specificity, and positive and negative predictive values, with their 95% confidence intervals, for smear and ligase chain reaction assay. The results were compared with the final clinical diagnosis using a positive culture as the gold standard to diagnose active tuberculosis. Positive ligase chain reaction results that were culture negative in patients with tuberculosis on treatment or in patients with old healed tuberculosis were felt to reflect the high sensitivity of the ligase chain reaction assay in detecting non-viable mycobacterial DNA, but were considered to be false positive in the final analysis.

**Results**

A total of 2420 specimens were tested in parallel by smear, ligase chain reaction, and culture assays over a 14 month period. Thirty nine patients had specimens which tested positive for smear (n=22), ligase chain reaction (n=37), or culture for M tuberculosis (n=33; table 1).

During this period there were two atypical mycobacteria cultured in the laboratory from sputum specimens (M malmoense and M chelonae). Both were smear and ligase chain reaction negative and were considered non-pathogenic.

In the group with newly diagnosed culture positive tuberculosis (n=32) there were 20 men (63%) and 12 women (37%) with a median age at diagnosis of 43 years (range 18–87). Nineteen of 28 patients on whom smoking information was available (68%) were current smokers; 29 patients had pulmonary tuberculosis (91%), two had renal tuberculosis (6%), and one had both pulmonary and renal tuberculosis (3%). Specimens with smear positive in 21 cases (66%), ligase chain reaction positive in 30 cases (94%), and culture positive in 32 cases (100%). In the two specimens which were ligase chain reaction negative but culture positive (one sputum, one pleural fluid) there were no recognisable similarities that might suggest a reason for the negative ligase chain reaction assay. In the smear positive group all specimens were ligase chain reaction and culture positive. In the smear negative group 18% of specimens were ligase chain reaction negative. Discrepant results between ligase chain reaction and culture were in the smear negative group.

In the patient with a relapse of tuberculosis sputum smear, ligase chain reaction, and culture were all positive. In the group with tuberculosis who were receiving treatment before ligase chain reaction testing (n=3) all specimens were sputum and were ligase chain reaction positive but smear and culture negative. Similarly, in the group with old healed tuberculosis (n=3) all specimens were sputum and were ligase chain reaction positive but smear and culture negative (median period elapsed since diagnosis was 2 years (range 2–10)).

In amalgamating the data, using a positive culture as the gold standard for the diagnosis of active tuberculosis, the Abbott ligase chain reaction M tuberculosis assay, when reported as positive, had a sensitivity of 93.9% (95% confidence interval (CI) 80 to 99), a specificity of 99.8% (95% CI 99.4 to 99.9), a positive

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The ligase chain reaction based assay for the detection of M tuberculosis complex was run according to the manufacturer’s instructions. This uses the nucleic acid amplification method ligase chain reaction to detect the presence of M tuberculosis DNA directly in clinical specimens. Sample preparation involves cell lysis to release DNA. Four oligonucleotide probes in the ligase chain reaction assay recognise and hybridise to a specific target sequence within the chromosomal DNA of the M tuberculosis complex. This nucleic acid sequence encodes for protein antigen b9, a gene sequence which appears to be specific to the complex strains examined to date.10 The oligonucleotides are designed to be complementary to the target sequence so that, in the presence of target, the probes will bind adjacent to one another. They can then be enzymatically joined to form the amplification product which subsequently serves as an additional target sequence during further rounds of amplification. Micropolypeptide enzyme immunoassay (MEIA) detects the product of the ligase chain reaction on the Abbott analyser. Culture was performed with Lowenstein-Jensen and pyruvate slopes incubated for 8 and 12 weeks for smear negative and smear positive samples, respectively.

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Ligase chain reaction in clinical practice

Table 2 Comparison of smear and ligase chain reaction in culture positive tuberculosis

<table>
<thead>
<tr>
<th></th>
<th>Smear</th>
<th>Ligase chain reaction</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sensitivity</td>
<td>66.7% (48.2 to 82.0)</td>
<td>93.9% (80 to 99)</td>
</tr>
<tr>
<td>Specificity</td>
<td>100% (99.8 to 100)</td>
<td>99.8% (99.4 to 99.9)</td>
</tr>
<tr>
<td>Positive predictive value</td>
<td>100% (84.6 to 100)</td>
<td>83.8% (68.0 to 93.8)</td>
</tr>
<tr>
<td>Negative predictive value</td>
<td>99.5% (99.2 to 99.8)</td>
<td>99.9% (99.7 to 99.9)</td>
</tr>
<tr>
<td>p value (χ² test)</td>
<td>&lt;0.0001</td>
<td>&lt;0.0001</td>
</tr>
</tbody>
</table>

Values are percentages with 95% confidence intervals in parentheses.

The χ² test had a two tailed p value of <0.0001.

Discussion

The data presented here suggest that the ligase chain reaction assay is a sensitive and specific tool for the detection of culture positive tuberculosis in clinical practice and may, in fact, be more sensitive and specific than other commercially available PCR based diagnostic tests for tuberculosis. However, this study was undertaken specifically to address the clinical efficacy of the ligase chain reaction assay in comparison with culture.

Although a strong presumptive diagnosis of pulmonary tuberculosis can be established by the finding of acid fast bacilli on microscopic examination in 50–60% of patients with this disease, identification of *M tuberculosis* in culture is required for confirmation. Using conventional solid media for culture, results are not available for 3–6 weeks. Conventionally, the smear positive patient is started on anti-tuberculous treatment immediately, but delay in initiating drug treatment in the smear negative patient may result in progression of the disease.

Although licensed only for respiratory specimens (sputum and bronchoalveolar lavage fluid), the ligase chain reaction assay is sensitive and specific in the diagnosis of extra-pulmonary tuberculosis. The assay was positive in two of three culture positive pleural fluid specimens and in both of two culture positive urine specimens in this trial.

Mycobacterial DNA may remain in the body many years after primary or post-primary infection, so the ligase chain reaction assay may be positive in clinical specimens from a patient with old (healed) tuberculosis and cannot determine therapeutic success or failure following antimicrobial treatment. As with any diagnostic test, results should be interpreted in the context of clinical findings. Its main advantage is speed, with results available within an eight hour working day. Although smear results are usually available within a similar period, the sensitivity of the ligase chain reaction assay is more comparable to that of culture than smear (table 2). In clinical practice this suggests a definite role for the ligase chain reaction assay in patients with negative smear but positive ligase chain reaction results in whom active *M tuberculosis* infection is clinically suspected. A useful role is also suggested in patients who have smear positive, ligase chain reaction negative specimens. Such results would suggest infection with mycobacteria other than tuberculosis (MOTT) as ligase chain reaction is specific for the *M tuberculosis* complex. There were no cases of *M bovis* in this series. This is consistent with the low prevalence of *M bovis* infection in humans in this region.

As two patients were ligase chain reaction negative and culture positive for *M tuberculosis* and six were ligase chain reaction positive but smear and culture negative (table 1), it is not anticipated that the ligase chain reaction assay will replace culture as a diagnostic tool for tuberculosis. Rather, the authors conclude that it is a useful primary screening test for tuberculosis, offering speed (8 hours vs 3–6 weeks) and discrimination (*M tuberculosis* vs MOTT) in the early stages of diagnosis, and complementing traditional smear and culture techniques.


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