Diagnostic potential of 16 kDa (HspX, α-crystalline) antigen for serodiagnosis of tuberculosis


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Background & objectives: Tuberculosis (TB) is a public health problem worldwide. Rapid and accurate diagnosis of tuberculosis is crucial to facilitate early treatment of infectious cases and to reduce its spread. The present study was aimed to evaluation of 16 kDa antigen as a serodiagnostic tool in pulmonary and extra-pulmonary tuberculosis patients in an effort to improve diagnostic algorithm for tuberculosis.

Methods: In this study, 200 serum samples were collected from smear positive and culture confirmed pulmonary tuberculosis patients, 30 tubercular pleural effusions and 21 tubercular meningitis (TBM) patients. Serum samples from 36 healthy, age matched controls (hospital staff), along with 60 patients with non-tubercular respiratory diseases were also collected and evaluated. Humoral response (both IgG and IgA) was looked for 16 kDa antigen using indirect ELISA.

Results: Sensitivity of detection in various categories of pulmonary TB patients ranged between 73.8 and 81.2 per cent. While in the extra-pulmonary TB samples the sensitivity was 42.8 per cent (TBM) and 63.3 per cent (tubercular pleural effusion). The test specificity in both the groups was high (94.7%). All of the non-disease controls were negative. Among non-tubercular disease controls, five patients gave a positive humoral response against 16 kDa.

Interpretation & conclusions: Serodiagnostic tests for TB have always had drawbacks of suboptimal sensitivity and specificity. The antigen used in this study gave encouraging results in pulmonary TB only, while in extra-pulmonary TB (tubercular meningitis and tubercular pleural effusion), this has shown a limited role in terms of sensitivity. Further work is required to validate its role in serodiagnosis of TB especially extra-pulmonary TB.

Key words ELISA - EPTB - serodiagnosis - tuberculosis - 16 kDa

Tuberculosis (TB) is a major public health problem worldwide and was responsible for 8.8 million incident cases of TB and 1.45 million deaths in the year 2010¹. The failure to diagnose TB accurately and rapidly is a key challenge in curbing the epidemic², ³. Rapid and accurate diagnosis of tuberculosis is crucial to facilitate early treatment of infectious cases and thus to reduce its spread.
TB diagnosis largely depends upon clinical examination and radiographic findings, mainly confirmed by sputum microscopy and bacterial culture\(^3\). Many alternative methodologies have been applied in TB diagnosis, such as PCR and cell-mediated immune response reactions\(^4\). These methods require trained personnel and specific laboratory conditions, which hinder their implementation in many areas of high TB endemicity.

Extra-pulmonary TB (EPTB) remains an important diagnostic and therapeutic problem. The diagnosis of extra-pulmonary tuberculosis is challenging due to the paucibacillary nature of the specimens, inability to access the site of disease activity, the lack of adequate sample volumes and the presence of inhibitors that undermine the performance of nucleic acid amplification-based techniques. Serological tests may prove useful with the advantages of speed, technical simplicity, possible adaptation towards point-of-care formats and low cost\(^5\).\(^6\).

The availability of numerous well characterized \textit{M. tuberculosis} proteins has revived interest in the serological diagnosis of tuberculosis. Several promising antigens have been reported such as 16 kDa, 45 kDa, antigen 85 complex (30 kDa), 65 kDa Hsp, 88 kDa, 38 kDa, ESAT-6, CFP-10, etc\(^8\). Antibody response to the 38 kDa in pulmonary TB has been extensively studied, and there are a few reports about the utility of the 16 kDa-based serological tests in pulmonary and extra-pulmonary TB\(^7\).\(^9\).

The aim of the present study was to explore the serodiagnostic potential of 16 kDa (HspX, Rv2031c, or \(\alpha\)-crystalline) antigen in different groups of tuberculosis patients. 16 kDa is an immunodominant antigen that is recognized by the majority of patients with active tuberculosis\(^10\). Production of Rv2031c appears to increase as the bacteria go into the metabolically resting stage and decreases as they revert to exponential growth\(^11\), thus providing a constant source of antigen in cultures and possibly \textit{in vivo}.

**Material & Methods**

A total of 347 subjects (age range= 18 to 65 yr, male= 235, female= 112) were recruited between June 2007 to July 2010 (Table I). All patients and control subjects gave informed consent prior to sampling. The cross-sectional study protocol was reviewed and approved by the ethics committee of the All India Institute of Medical Sciences (AIIMS), New Delhi. Detailed clinical history and findings were recorded in a case record form.

A total of 200 pulmonary TB (PTB) patients were enrolled prospectively from Comprehensive Rural Health Services Project (CRHSP), Centre for Community Medicine, Ballabgarh, Haryana (n=120) and Rajan Babu Institute for Pulmonary Medicine & Tuberculosis (RBIPMT), New Delhi (n=80); and were classified 78 category I PTB patients (new sputum smear positive), 42 category II PTB patients (sputum smear positive after default or relapse/treatment failure) and 80 multi-drug resistant PTB patients.

Sputum (5-10 ml) and blood samples (2-3 ml) were collected within two weeks of initiation of anti-tuberculosis treatment (ATT) from category I and category II patients (from CRHSP). MDR PTB patients admitted at RBIPMT were on ATT at the time of inclusion in the study. All the patients included in the study had the sputum culture positive for \textit{M. tuberculosis}. After collection, samples were transported in cold chain.

Twenty one in-patients with tuberculous meningitis (TBM) were enrolled from the Department of Neurology, AIIMS, New Delhi. Sixteen patients were clinically and radiologically confirmed\(^12\) and cerebrospinal fluid (CSF) of five patients grew \textit{M. tuberculosis} on culture. Blood sample (2-3 ml) was collected within two weeks of initiation of ATT.

Thirty patients with tubercular pleural effusion were enrolled from CRHSP, Ballabgarh, Haryana and AIIMS, New Delhi. Patients were clinically and radiologically confirmed as the microbiological confirmation could not be obtained for all patients. A total of 60 subjects, who were suffering from respiratory/lung diseases (other than tuberculosis) were included as disease controls from AIIMS, New Delhi. Lung cancer (n=21), asthma (n=9), pneumonia (n=7), chronic obstructive pulmonary disorder (n=3), interstitial lung disease (n=3), sarcoidosis (n=3), respiratory distress (n=3), allergic broncho pulmonary aspergillosis (n=3), bronchiectasis (n=2), occupational lung disease (n=1), Wegener’s granulomatosis (n=1), bronchiolitis obliterans (n=1), metabolic encephalopathy (n=1) formed the spectrum. Sputum/induced sputum samples were obtained. All above disease controls were evaluated bacteriologically for \textit{M. tuberculosis} using culture and AFB smear and found negative. Forty six subjects had \textit{M. bovis} BCG vaccination history. The purified protein derivative (PPD) status of the patients was unknown.
Thirty six healthy individuals (with no known history of active TB) were enrolled from the staff working in Department of Microbiology, AIIMS, New Delhi, as healthy controls or non-disease controls. All these were healthy with no apparent signs of any disease. All of them had BCG vaccination history. The purified protein derivative (PPD) status of the subjects was not known.

Sample processing: Sputum samples were processed using standard NALC-NaOH method\textsuperscript{13} and smears were examined after Ziehl–Neelsen staining. Processed samples were inoculated in MGIT (Mycobacterial growth indicator tube) 960 non-radiometric automated isolation system (BD, USA). MGIT tube was supplemented with 0.5 ml of oleic acid-albumin-dextrose-catalase along with mixture of 5 antibiotics (PANTA i.e., polymyxin B, amphotericin B, nalidixic acid, trimethoprim and azlocillin) and 0.5 ml of decontaminated sample. 

\textit{M. tuberculosis} complex and non-tuberculous mycobacteria were differentiated using p-nitrobenzoic acid (PNBA) test (as recommended in MGIT-960 protocol). Drug susceptibility testing (DST) was performed using 1 per cent proportion method using above automated culture system to reassure the MDR status (as recommended in MGIT-960 protocol). Serum samples were separated and immediately stored in -80°C till further use.

**ELISA protocol:** ELISA was performed with some modifications\textsuperscript{14,15} to estimate the humoral responses (IgG and IgA, in the same well) against native 16 kDa antigen. Briefly, the 96-well microtiter plates (Maxisorp Nunc, Denmark) were coated with native antigen in PBS (50 µl/well of 5 µg/ml) overnight at 4°C. Next day, plates were washed three times with PBS-T, blocked with 1 per cent BSA in PBS-T (blocking buffer) for two hours at 37°C and followed by three washings. Subsequently, 100 µl of diluted sera (1:100 in 0.1 x blocking buffer) from patients and healthy subjects was added and the plates were incubated for 90 min at 37°C. After six washes with PBS-Tween 20 (0.05% Tween 20 in PBS), a mixture of alkaline phosphatase-conjugated protein A (1:2,000) and anti-human immunoglobulin A (IgA; 1:1,000) (Sigma-Aldrich, St. Louis, USA) was added to each well and the plates were incubated for 1 h at 37°C. The wells were washed six times with PBS-T, and the bound enzyme-conjugated antibodies were detected with p-nitrophenylphosphate substrate (Sigma, USA), (1 mg/ml p-nitrophenylphosphate in 10% diethanolamine buffer containing 0.5 mM MgCl$_2$, pH 9.8). The plates were read at 405 nm in ELISA plate reader (Microscan, ECIL, India).

Data analysis: The cut-off was determined by calculating the mean optical density (OD) obtained with sera from healthy individuals plus 3 standard deviations.

<table>
<thead>
<tr>
<th>Sr. no.</th>
<th>Subjects</th>
<th>Total no.</th>
<th>Male/ female</th>
<th>Mean age (yr) ± SD</th>
<th>Inclusion criteria</th>
<th>Positive for 16 kDa antigen</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.</td>
<td>Category I pulmonary TB</td>
<td>78</td>
<td>54/24</td>
<td>31.1 ± 11.9</td>
<td>Culture confirmed for \textit{M. tuberculosis}</td>
<td>63</td>
</tr>
<tr>
<td>2.</td>
<td>Category II pulmonary TB</td>
<td>42</td>
<td>30/12</td>
<td>35.5 ± 12.4</td>
<td>Culture confirmed for \textit{M. tuberculosis}</td>
<td>31</td>
</tr>
<tr>
<td>3.</td>
<td>MDR pulmonary TB</td>
<td>80</td>
<td>47/33</td>
<td>33.2 ± 9.8</td>
<td>Culture confirmed for \textit{M. tuberculosis}, Sensitivity confirmed MDR-TB</td>
<td>65</td>
</tr>
<tr>
<td>4.</td>
<td>Tuberculous meningitis (TBM)</td>
<td>21</td>
<td>13/8</td>
<td>34.3 ± 14.6</td>
<td>Clinically &amp; radiologically confirmed (five samples bacteriologically confirmed)</td>
<td>9</td>
</tr>
<tr>
<td>5.</td>
<td>Tubercular pleural effusion</td>
<td>30</td>
<td>21/9</td>
<td>37.3 ± 14.9</td>
<td>Clinically &amp; radiologically confirmed</td>
<td>19</td>
</tr>
<tr>
<td>6.</td>
<td>Non-disease controls</td>
<td>36</td>
<td>27/9</td>
<td>31.2 ± 6.9</td>
<td>Healthy individuals with no apparent signs for any disease</td>
<td>0</td>
</tr>
<tr>
<td>7.</td>
<td>Disease controls (non-tubercular)</td>
<td>60</td>
<td>43/17</td>
<td>45 ± 13.1</td>
<td>Disease confirmed</td>
<td>5*</td>
</tr>
</tbody>
</table>

*One each of lung cancer, pneumonia, ILD, sarcoidosis, respiratory distress
deviations (SD). Each assay was repeated three times. Two or three positives out of three ELISAs were considered as positive. Sensitivity and specificity were calculated by standard methods. Scatter plot and ROC (receiver operative characteristic) curves were plotted using GraphPad Prism software, version 5, USA. ROC curve describes probabilities of correct and incorrect results at different cut-off values. The area under ROC curve reflects the accuracy of a test16.

Results

The 347 subjects when grouped into different categories, 78 were in category I PTB, 42 were in category II PTB, 80 were MDR PTB, 21 were tuberculous meningitis, 30 were pleural effusion, 36 were non-disease controls and 60 were disease controls (other than tuberculosis) (Table I).

Fig. 1 shows the scatter plot of antibody response in different groups of patients and controls. Each dot represents the ΔO.D. (ΔO.D.= O.D. of sample - cut-off value) for individual patient. All dots above X-axis are positive O.Ds. All dots below X-axis are negative O.Ds.

16 kDa antigen was used for detection of humoral responses (IgG and IgA) in these subjects using ELISA and sensitivity, specificity and other parameters were predicted (Table II). The diagnostic sensitivity of 16 kDa antigen with category I PTB, category II PTB and MDR PTB was found to be 80.7, 73.8 and 81.2 per cent, respectively. The sensitivity was found to be comparatively lower in category II PTB than the category I PTB and MDR PTB. While in the extra-pulmonary TB i.e. tuberculosis meningitis and tubercular pleural effusion cases, the diagnostic sensitivity was 42.8 and 63.3 per cent, respectively. Specificity was high (>94%) in PTB as well as extra-pulmonary tuberculosis groups (Table II).

None of the non-disease controls was positive for antibodies to 16 kDa, however, five disease controls were found to be positive. These patients were being treated for lung cancer (1), sarcoidosis (1), interstitial

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**Table II. Sensitivity, specificity and likelihood ratios of 16 kDa antigen ELISA for antibody detection**

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>Category I PTB (n=78)</th>
<th>Category II PTB (n=42)</th>
<th>MDR PTB (n=80)</th>
<th>Tuberculous meningitis (n=21)</th>
<th>Tubercular pleural effusion (n=30)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sensitivity (%)</td>
<td>80.7 (69.9 to 88.4)</td>
<td>73.8 (57.6 to 85.6)</td>
<td>81.2 (70 to 88.7)</td>
<td>42.8 (22.5 to 65.5)</td>
<td>63.3 (43.9 to 79.4)</td>
</tr>
<tr>
<td>Specificity (%)</td>
<td>94.7 (87.7 to 98)</td>
<td>94.7 (87.7 to 98)</td>
<td>94.7 (87.7 to 98)</td>
<td>94.7 (87.7 to 98)</td>
<td>94.7 (87.7 to 98)</td>
</tr>
<tr>
<td>Likelihood ratio for positive test result (LR+)</td>
<td>15.5</td>
<td>14.2</td>
<td>15.6</td>
<td>8.2</td>
<td>12.2</td>
</tr>
<tr>
<td>Likelihood ratio for negative test result (LR-)</td>
<td>0.20</td>
<td>0.27</td>
<td>0.19</td>
<td>0.60</td>
<td>0.38</td>
</tr>
</tbody>
</table>

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**Fig. 1.** (I) Scatter plot of humoral responses in serum samples of non-disease controls, disease controls, category I PTB, category II PTB, MDR-TB, tuberculous meningitis (TBM) and tubercular pleural effusion patients. Each dot represents the ΔO.D. (ΔO.D. = O.D. of sample – cut-off value) for individual patient. All dots above X-axis are positive O.Ds. All dots below X-axis are negative O.Ds.
lung disease (ILD 1), pneumonia (1) and respiratory distress (1).

Discussion

In the present study, we evaluated the 16 kDa antigen, in PTB and EPTB (TBM and tubercular pleural effusion) patients, and compared the antibody assay with other diagnostic modalities. In our study, none of the non-disease controls was positive for antibodies to 16 kDa. In a TB endemic country like India a majority of the population is likely to be latently infected or BCG-vaccinated; the absence of any humoral response against mycobacteria gives credence to high specificity of the antigen for detecting active disease. Geluk et al. have shown most M. tuberculosis infected or exposed individuals responded well to HspX, whereas significantly lower responses were observed in unexposed individuals, including BCG-vaccinated individuals. Rabahi et al. have shown higher HspX in the M. tuberculosis exposed individual and good indicator for new infection. Davidow et al. found higher humoral response among inactive TB than the active TB.

A meta-analysis performed by Steingart et al. has reported high specificity for detection of pulmonary tuberculosis (86 to 100%), using the commercially available “pathozyme TB complex plus” (ELISA based antibody detection, utilizes the recombinant 38 kDa and 16 kDa) in various studies. Another study has shown 98 per cent specificity in a large number of Polish population using the above test. Raja et al. have shown a specificity of 94 per cent using 16 kDa antigen for antibody detection in pulmonary tuberculosis patients and controls including non-tuberculous lung disease and healthy subjects.

Several antigens have been shown to have strong immunodiagnostic potential. A couple of studies are available for the 16 kDa molecule, a polypeptide belonging to the α-crystallin family of low molecular weight heat shock proteins. The coding gene (Rv2031c) for this antigen has been found exclusively in the M. tuberculosis complex as shown by DNA hybridization studies. It has been reported as a dominant protein, produced in the static growth phase or under oxygen deprivation and is essential for bacterial replication inside macrophages.

In non-tubercular disease controls, five of 60 controls gave the non-specific positive humoral response. Though sputum/induced sputum were obtained from above disease control group, all were negative for AFB smear microscopy and MGIT-960 rapid culture. All the care had been employed at the time of enrollment of above disease controls to rule out history of active TB. An earlier study has shown that antibodies against 16 kDa do not cross-react with common environmental mycobacteria. Julian et al. have shown some non-specificity in pneumonia.

![Fig. 2. Receiver operative characteristic (ROC) curve for (A): PTB (category I PTB, category II PTB and MDR). Area under ROC curve 0.9083, 95% CI (confidence interval) 0.8739 to 0.9428, Std. error 0.01758, P<0.001. Small bars on the ROC curve showing the cut-offs. (B): EPTB (tuberculous meningitis and tubercular pleural effusion) area under ROC curve: 0.7571, 95% CI 0.6589 to 0.8554, Std. error 0.05013, P<0.001.](image-url)
population using 16 kDa antibody detection, antibodies to 16 kDa have been shown among non-TB lung diseases such as asthma, lung cancer. The loss of specificity may be due to latent infection or to nonspecific hyperglobulinaemia, common in bronchial diseases.

Our results have shown a significant (P<0.05) higher antibody response in PTB than EPTB for 16 kDa antigen. Many of the serological studies have been reported from patients with extensive pulmonary disease. The group that would most benefit from serological tests is the one with smear negative or extra-pulmonary TB. In this group of patients, the sensitivity of different assays was reported to be significantly lower than in the confirmed cases. The extensive extra-cellular bacterial replication in cavities and the continued shedding of antigens is possibly enhanced in the pulmonary cavitary environment. The paucibacillary and often walled off EPTB lesions may be different in producing lesser amount of antigen and hence may take longer to generate an antibody response. The study patients were sampled within a few days of the clinical diagnosis and hence may not have shown an adequate serological response.

The sensitivity was much higher in category I, category II PTB and MDR PTB patients. Repeated immune stimulation by mycobacterial antigens and higher antigenic load because of multiplying mycobacteria in MDR-PTB and category II PTB may explain this.

Serodiagnostic tests for tuberculosis have never been very successful due to suboptimal sensitivity and specificity. To discourage the rampant use of commercial TB serodiagnostic tests, WHO has issued a policy note discouraging the use of current serodiagnostic tests.

The present study has limitations such as the serologic tests were performed retrospectively; using serum that had been stored at -80°C, although no serum had been thawed more than once. The total number of EPTB (TBM and tubercular pleural effusion) subjects enrolled in the study was small. The results in our study were encouraging only in PTB group; and in tuberculous meningitis and tubercular pleural effusion patients were not up to expectations. The use of additional antigen(s) combinations or epitopes need to be evaluated for serodiagnosis.

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References


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