ORIGINAL ARTICLE

Single-step QuantiFERON screening of adult contacts: a prospective cohort study of tuberculosis risk

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ABSTRACT

Background The effectiveness of tuberculosis (TB) contact screening programmes using interferon γ release assays remains uncertain as prospective contact TB risk is not well characterised.

Objectives To quantify 2-year TB risk and evaluate screening performance with single-step QuantiFERON TB Gold-In Tube (QFT) in adult contacts. To compare TB risk between QFT tested subgroups stratified by exposure type (smear positive pulmonary (SP) versus non-smear positive (NSP) TB) and age (younger (16–35 years) versus older (≥36 years)).

Methods Screening involved QFT testing in older contacts of SP and all younger contacts, 8–12 weeks after index notification. Chemoprevention (3RH) was offered to QFT positive (+) younger adults. TB risk was determined in a prospective cohort study.

Results 43 TB events occurred in 1769 adult contacts observed for median 717 days (2-year rate (95% CI)=2.5% (1.7 to 3.2)). Index-contact strain matching was demonstrable for 18 of 22 (82%) paired samples. No contacts (0/98) receiving 3RH developed TB. 215 of 817 appropriately tested adults (26.3%) were QFT+. 14 of 112 untreated QFT+ adults developed TB (2-year rate (95% CI)=13.4% (7.7 to 21.1)). The model required 35 contacts screened with QFT to identify one contact developing TB at 2 years. TB rates were comparable in QFT + contacts of SP and NSP (rate ratio (RR)=0.98, p=0.962). For QFT+ older contacts, the disease rate was lower (8.9% (3.3 to 19.1)) and similar to the overall group rate (RR=1.4, p=0.503).

Conclusions QFT based single-step contact screening is effective in young adults.

INTRODUCTION

Preventing future tuberculosis (TB) requires screening programmes to identify and treat latent infection with Mycobacterium tuberculosis (LTBI) in high risk populations. However, LTBI is associated with a variable future risk of active TB that is complex, incompletely understood and poorly characterised.1 2 Whether greater biological specificity of interferon γ (IFNγ) release assays (IGRAs), compared with the traditional tuberculin skin test, is associated with identification of LTBI at a higher prospective risk of TB remains a subject of considerable debate and the focus of two recent meta-analyses.3 4 However, discordance in outcomes and conclusions between the meta-analyses is striking and attributable to differences in study selection that significantly influence the context of LTBI study. The analysis of Rangaka et al3 was confined to studies performed in low and middle income countries with a moderate to high TB burden. In contrast, Diel et al3 analysed a smaller number of studies from high income, low TB prevalence settings. Even for the two studies performed in a low prevalence setting that have specifically evaluated TB risk with IGRAs, the focus differs between them in reported prospective 2-year TB risk.5 6 Important differences in study design, objectives and population characteristics are likely to have contributed to this difference.
A clear need therefore exists for further studies to better inform IGRA based screening policies.\(^7\) \(^8\) Since January 2007, we have performed QuantiFERON TB Gold-In Tube (QFT) based single-step adult contact screening at our centre that is adapted from 2006 UK National Institute for Health and Clinical Excellence guidance.\(^9\) All contacts were followed prospectively for occurrence of secondary TB events. Here we present the findings of a longitudinal cohort analysis that quantifies 2-year TB risk in contacts of both SP and other non-smear positive (NSP) TB. We examine whether prospective TB risk with QFT defined LTBI differs between recent adult contacts according to their type of exposure and age and present indices of effectiveness with a single-step model of screening with QFT in clinical practice.

**METHODS**

**Study population**

The study population included all notified cases of active TB and associated asymptomatic contacts entering the screening programme between 1 January 2007 and 5 June 2009. Cases of active TB in the contact population were recorded until 15 March 2010.

**Active TB**

A diagnosis of active TB was made in accordance with accepted criteria.\(^9\) IGRA testing is not included in the diagnostic evaluation for active TB in adults at our centre. For cases with culture confirmation, genotyping of the isolate using PCR based identification of variable number of tandem repeats for mycobacterial interspersed repetitive units (VNTR-MIRU) of 15 discriminatory loci\(^10\) was performed at the HPA Regional Mycobacteriology Laboratory in Birmingham.

**Screening algorithm**

We designed and employed a single-step model of screening for adult contacts (age ≥ 16 years) that was adapted from 2006 UK National Institute for Health and Clinical Excellence guidance\(^9\) (figure 1). In brief, contact tracing procedures were initiated immediately after index notification and screening for active TB performed using a symptom questionnaire. QFT testing was performed according to the manufacturer’s instructions (Cellestis, a QIAGEN company, Hilden, Germany) in all contacts under 36 years and in contacts of any age exposed to SP TB after 8–12 weeks. All QFT positive adult contacts under 36 years were offered chemoprophylaxis with rifampicin and isoniazid for 3 months.
(3RH), unless drug resistance was identified in the source case. QFT positive contacts aged ≥56 years and younger QFT positive contacts who declined chemoprophylaxis had active follow-up with clinical assessment and a chest radiograph at 6-monthly intervals for 2 years after index notification. QFT negative adults and untested contacts aged ≥56 years were followed-up once after 3 months. Passive follow-up is maintained for all contacts who remained in the region through a centralised system of rapid access and our local networked electronic TB database (TBIT) (online supplement).

**Screening timeline and categorisation of contact TB**

By definition, contact TB risk was modifiable with screening if an opportunity for chemoprevention existed prior to disease onset (figure 1). On this basis, contact TB identified ≤14 weeks after index notification (early and co-prevalent TB) was not preventable and TB occurring after 14 weeks (late disease) was potentially modifiable with screening (online supplement).

**Statistical methodology**

The primary outcome of this study was cumulative 2-year risk of active TB in QFT positive adult contacts of SP and NSP TB, using our screening algorithm. NSP TB included all non-pulmonary cases and smear negative pulmonary TB. Secondary outcomes were quantification of performance with QFT as a single-step screening tool for targeted delivery of chemoprophylaxis; comparison of TB rates between younger (<56 years) and older adults; and characterisation of factors associated with failure of the screening model to prevent contact TB. All contact cases were included for characterisation of factors associated with screening failure. Contacts with protocol deviation (inappropriate QFT testing and/or inappropriate chemoprophylaxis) and QFT tested contacts diagnosed within 2 weeks of testing (QFT result representative of active TB and not LTBI) were excluded from QFT based longitudinal analyses.

Between-group statistical comparisons were performed using the independent t test (parametric variables) and Mann–Whitney test (non-parametric variables). Kaplan Meier analysis was performed to estimate TB risk in the prespecified contact groups and compared between them using the log-rank test. The number needed to screen (NNS) and number needed to treat (NNT) were calculated (online supplement) as respective indices of screening and therapy effectiveness with our model.

Performance measures of QFT as a screening tool included population independent (sensitivity, specificity and positive and negative likelihood ratios) and population dependent (positive predictive value and negative predictive value; online supplement) parameters. These were computed with contingency tables for all contacts and separately in contacts of SP and NSP TB. Contingency tables were also used to calculate TB rate ratios. The 95% CIs were determined using Wald’s method and significance testing performed with the χ² test.

Forward stepwise logistic regression was performed (online supplement) to model the association of prespecified variables (see online supplementary table S2) with the QFT result in all tested contacts.

SPSS V16 (SPSS, Inc.) was used to perform survival analyses, and significance testing with the log rank test and logistic regression. Prism V5 (GraphPad Software, Inc., California, USA) was used to perform analyses of contingency tables. For all analyses, the significance threshold was <0.05.

The study was approved by the Leicestershire TB Management Board; the local research ethics committee was consulted and formal ethical approval was not required. No external funding was obtained.

**RESULTS**

**Study population and uptake of the screening programme**

Close contacts were identified for 505 of 628 (80%) notified TB cases (see online supplementary table S1). Overall, 1769 of 2401 contacts (74%) were aged above 15 years and included in the study (figure 2). The overall culture confirmation rate in the cohort with index TB was 60% and significantly higher for pulmonary TB, compared with non-pulmonary disease (76% vs 41%; p<0.001).

In 34 adults (1.9%) protocol deviations were identified, and comprised inappropriate screening and treatment of older adults (figure 2). A total of 817 of 1100 eligible adults (74.4%) were QFT tested. The mean (±SD) time to testing after index notification was 95±22 days. The proportion of eligible contacts tested did not differ across prespecified subgroups, stratified by type of disease exposure or age. In all, 98 of 147 (66.0%) eligible QFT positive contacts accepted chemoprophylaxis and treatment completion confirmed in 76 (74.6%) of this group. In all treated subjects, chemoprophylaxis was well tolerated with no adverse events requiring cessation of therapy.

**Outcome of QuantiFERON testing**

Overall, 215 (26.3%) of the 817 appropriately tested contacts were QFT positive (figure 2). An indeterminate result occurred for one contact. Five QFT positive contacts were diagnosed with TB at the time of screening (ie, within 2 weeks of testing). Modelling identified the contact’s age, place of birth (UK or abroad) and disease type of the index case (SP or NSP) to be independently associated with the QFT result. Of these, index disease type was most strongly associated with an OR (95% CI) for a positive QFT of 5.6 (3.4–9.1) for contact with SP TB, compared with NSP disease (see online supplementary table S2). Overall, the proportion of contacts who were QFT positive was 20.2% for non-pulmonary, 18.6% for smear negative pulmonary and 54.8% for SP TB, respectively.

**Secondary TB in contacts**

Contacts were observed prospectively for a median of 717 days (range 283–1157 days). Based on published migration figures (online supplement),12 we estimated 95 contacts would have been lost to follow-up, with a risk of TB occurring in less than two contacts over the median period of observation. In all, 43 cases of active TB were identified in contacts of 37 index cases. A table summarising details for each contact case is provided (see online supplementary table S3). One of 35 tested adults was HIV positive at the time of notification. A positive culture for *M tuberculosis* was obtained in 26 contact cases. For the remainder, TB was diagnosed on the basis of compatible clinical symptoms, signs and radiological features coupled with either clinical and radiological response following a full course of anti-tuberculous therapy alone (N=9) or additionally with histological evidence of necrotising granulomatous inflammation (N=8) (see online supplementary table S2). Genotype data for the isolate were available for 22 pairs of index cases and associated contact cases and genotypically matched in 18 pairs (82%). There were no differences in characteristics between subgroups with matched and unmatched results.

The median time to contact TB after notification of the index case was 171 days (IQR 95–287 days). Co-prevalent TB was diagnosed in one contact after 16 days and nine further contacts were categorised with early disease. For the remaining
33 contacts (76.7%) with late disease, three primary factors associated with failure to prevent TB were identified (online supplementary figure 1): non-adherence with the screening model (N=12); ineligibility for chemoprophylaxis due to age (N=15); and QFT negative at time of screening (N=6).

Testing with QuantiFERON and TB risk in contacts
For untreated contacts, the overall 2-year TB rate was 2.5% (95% CI 1.7 to 3.2) (table 1). Contacts of SP TB had a 4.8-fold higher risk of TB than contacts of NSP TB (p<0.001; figure S). There was no difference in the rate of TB between older and younger contacts (rate ratio (95% CI))=1.1 (0.6 to 2.1), p=0.678).

A total of 41 of 1769 contacts (2.3%) were excluded from QFT based analyses of TB risk and screening performance. This subgroup included eight contact TB events of whom one had co-prevalent disease, five were diagnosed at the time of QFT testing and two were cases that occurred in inappropriately screened contacts (figure 2). Summary characteristics of the cohort included are summarised in table 2. For untreated QFT positive contacts, the 2-year TB risk (95% CI) was 13.4% (7.7 to 21.1) (table 1) and did not differ between contacts of SP and NSP disease (figure S). There was no difference in the IFNγ response to mycobacterial antigens between QFT positive contacts of SP and NSP disease (figure S).
response of QFT positive contacts who did and did not develop TB, either in the proportion with an IFNγ titre >10 IU/ml (15.9% for non-progressors and 15.4% for progressors to TB; p=0.962) or in the IFNγ titre when this was below 10 IU/ml (1.1-fold difference, 95% CI 0.6 to 1.9; p=0.762).

Overall, QFT had a sensitivity of 70% and specificity of 85.9% for identifying LTBI progressing to TB at 2 years in our study cohort, using the single-step model. These and other indices of screening performance were comparable between untreated contact subgroups of SP and NSP TB (table 1). A positive QFT result informed a significantly higher 2-year TB risk in all prespecified subgroups except older contacts of SP disease. For this population, there was no difference between the overall disease rate and the rate in the QFT positive subgroup (rate ratio=1.4 (95% CI 0.5 to 5.9), p=0.503; table 1).

Based on the prevalence of QFT defined LTBI in our population, we estimated a NNS of 55 (95% CI 22 to 95) contacts with our model to identify one adult developing TB over 2 years, without chemopreventive therapy (table 1).

Six cases arose in contacts who were QFT negative at the time of screening, giving QFT a negative predictive value of 99% (95% CI 97.8 to 99.6) (table 1). Three out of four cases retested at the time of presentation with active disease were IGRA positive (two QFT, one T-SPOT.TB). All six cases were HIV seronegative. A positive culture was obtained for two cases without chemoprophylaxis.

DISCUSSION

In this study, we present longitudinal outcomes with a single-step QFT based programme of screening for adult contacts of TB. We have quantified 2-year TB risk for QFT positive LTBI in contacts of both SP and NSP TB and show that while a fivefold difference exists in the background rate of TB between the two cohorts, there is no difference in the TB rate between subgroups with QFT defined LTBI. We also demonstrate that the positive predictive value of QFT is lower in older contacts of SP TB and may not usefully risk stratify this group for targeted chemoprophylaxis. Our 2-year case rate of 13.4% was comparable with the rate reported by Diel et al and considerably higher than the figure of

![Figure 3](http://image-url.com)

**Figure 3** Overview of prospective 2-year tuberculosis (TB) risk in untreated adult contacts, stratified by exposure to smear positive pulmonary (SP) and non-smear positive (NSP) TB. The survival curves for TB occurrence are presented for all contacts of NSP and SP TB and for the subpopulations of each who were QuantiFERON TB Gold-In-Tube (QFT) tested and positive. The number of cases and size of the cohort from which they arise are presented with the key: Cases/Size of cohort. RR=2-year rate ratio (95% CI) of disease calculated between exposure subgroups (SP to NSP), as indicated. This figure is only reproduced in colour in the online version.

### Table 1 Two-year tuberculosis risk in untreated adult contact subgroups and indices of performance with QFT for screening

<table>
<thead>
<tr>
<th>Subgroups of contact exposure</th>
<th>All adult contacts</th>
<th>Non-smear positive</th>
<th>Smear positive (all)</th>
<th>Smear positive (≥36 years)</th>
</tr>
</thead>
<tbody>
<tr>
<td>2-year risk (cases/size of cohort)†</td>
<td>1.3 (0.8 to 2.1) [16/1263]</td>
<td>6.2 (4.6 to 8.1) [23/406]</td>
<td>6.5 (3.4 to 10.5) [12/211]</td>
<td></td>
</tr>
<tr>
<td>QFT positive</td>
<td>13.4 (7.7 to 21.1) [14/112]</td>
<td>13.4 (6.8 to 22.8) [10/80]</td>
<td>2.0 (0.9 to 4.2) [0.04]</td>
<td></td>
</tr>
<tr>
<td>QFT negative</td>
<td>1.1 (0.4 to 2.3) [6/601]</td>
<td>2.0 (0.8 to 5.4) [5/230]</td>
<td>4.8 (2.6 to 8.1) p=0.01</td>
<td></td>
</tr>
<tr>
<td>Rate ratio (QFT pos: All)‡</td>
<td>4.9 (2.7 to 8.7) p&lt;0.001</td>
<td>8.9 (3.1 to 25) p&lt;0.001</td>
<td>2.1 (1.0 to 4.2) p=0.041</td>
<td></td>
</tr>
<tr>
<td>Prevalence QFT positive/%</td>
<td>26.3 (23.2 to 29.0)</td>
<td>19.5 (16.0 to 23.0)</td>
<td>34.5 (29.6 to 39.4)</td>
<td></td>
</tr>
<tr>
<td>Sensitivity§</td>
<td>70.0 (45.7 to 88.1)</td>
<td>80.0 (28.4 to 99.5)</td>
<td>66.7 (38.4 to 88.2)</td>
<td></td>
</tr>
<tr>
<td>Specificity§</td>
<td>85.9 (83.0 to 88.4)</td>
<td>93.0 (89.9 to 95.3)</td>
<td>76.3 (71.0 to 81.0)</td>
<td></td>
</tr>
<tr>
<td>PPV§</td>
<td>12.5 (7.0 to 20.1)</td>
<td>12.5 (3.5 to 29.0)</td>
<td>12.5 (6.2 to 21.8)</td>
<td></td>
</tr>
<tr>
<td>NPV§</td>
<td>99 (97.8 to 99.6)</td>
<td>99.7 (98.5 to 99.9)</td>
<td>97.8 (95.0 to 99.3)</td>
<td></td>
</tr>
<tr>
<td>LR++</td>
<td>1.1 (0.4 to 2.3) [6/601]</td>
<td>0.3 (0.01 to 1.7) [1/371]</td>
<td>2.4 (0.8 to 5.4) [5/230]</td>
<td></td>
</tr>
<tr>
<td>Number needed to screen</td>
<td>35.0 (21.5 to 93.1)</td>
<td>41.5 (20.0 to 432)</td>
<td>40.0* (7.6 to 22.5)</td>
<td></td>
</tr>
<tr>
<td>Number needed to treat</td>
<td>6.8 (3.8 to 9.9)</td>
<td>6.5 (3.6 to 33.8)</td>
<td>7.0 (4.5 to 15.0)</td>
<td></td>
</tr>
</tbody>
</table>

All figures in parentheses are 95% CI of the mean unless stated otherwise.

†Two-year risk is presented as a cumulative percentage (95% CI). Figures in square brackets refer to the actual numbers of cases and the size of the cohort from which they arise, for each subgroup.

‡For rate ratio calculations, all contacts refers to the group selected for longitudinal analysis that was adherent with the screening algorithm (see Methods section).

§PPV, NPV, sensitivity and specificity are all presented as % values.

**p** Value >0.05.

<table>
<thead>
<tr>
<th>Year</th>
<th>Rate ratio (QFT pos: All)‡</th>
<th>Prevalence QFT positive/%</th>
<th>Sensitivity§</th>
<th>Specificity§</th>
<th>PPV§</th>
<th>NPV§</th>
</tr>
</thead>
<tbody>
<tr>
<td>2009</td>
<td>4.9 (2.7 to 8.7) p&lt;0.001</td>
<td>26.3 (23.2 to 29.0)</td>
<td>70.0 (45.7 to 88.1)</td>
<td>85.9 (83.0 to 88.4)</td>
<td>12.5 (7.0 to 20.1)</td>
<td>99 (97.8 to 99.6)</td>
</tr>
<tr>
<td>2010</td>
<td>8.9 (3.1 to 25) p&lt;0.001</td>
<td>19.5 (16.0 to 23.0)</td>
<td>80.0 (28.4 to 99.5)</td>
<td>93.0 (89.9 to 95.3)</td>
<td>12.5 (3.5 to 29.0)</td>
<td>99.7 (98.5 to 99.9)</td>
</tr>
<tr>
<td>2011</td>
<td>2.1 (1.0 to 4.2) p=0.041</td>
<td>34.5 (29.6 to 39.4)</td>
<td>66.7 (38.4 to 88.2)</td>
<td>76.3 (71.0 to 81.0)</td>
<td>12.5 (6.2 to 21.8)</td>
<td>97.8 (95.0 to 99.3)</td>
</tr>
<tr>
<td>2012</td>
<td>1.4* (0.5 to 3.9) p=0.503</td>
<td>39.1 (31.5 to 46.7)</td>
<td>62.5 (30.6 to 86.3)</td>
<td>62.5 (54.6 to 69.8)</td>
<td>8.1 (2.7 to 17.8)</td>
<td>96.9 (91.3 to 99.4)</td>
</tr>
</tbody>
</table>

All untreated 2.5 (1.7 to 3.2) [39/1669] 0.35 (0.18 to 0.68) 0.22* (0.04 to 1.2) 0.44 (0.21 to 0.90) 2.1* (0.24 to 1.5) 0.6* (0.24 to 1.5)

All contacts Non-smear positive Smear positive (all) Smear positive (≥36 years)

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3.1% reported by Kik et al.6 However, the latter included only immigrant contacts and the low positive predictive value of IGRA s supported their view that these assays discriminate poorly between recently acquired LTBI (at higher risk of secondary TB) and remote infection.13 In our study, older contacts were more likely to have remotely acquired LTBI, as evidenced by a greater proportion who were foreign born (table 2) and an independent association of increasing age with a positive QFT (see online supplementary table S2). Our finding that a positive QFT result was not significantly discriminatory for this group is therefore consistent with Kik et al and favours the view that QFT is less effective as a screening tool in populations with a significant burden of remote infection. Although limitations of QFT support the practice of age-limited screening,9 15 of 43 contact cases (35%) in our cohort developed TB, indicating that the retained positive pre-
screening and provision of early chemotherapy to QFT positive contacts, followed by a second phase of treatment after 6–12 weeks in QFT negative contacts, is possible that a proportion of early cases represented asymptomatic co-prevalent disease that would be unpreventable with any screening.

Table 2 Characteristics of adult contacts and subgroups included for QFT based longitudinal analyses

<table>
<thead>
<tr>
<th>Gender</th>
<th>All adult contacts (N=1769)</th>
<th>Contacts for QFT based longitudinal analysis (N=1728)</th>
<th>Sig (p value)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Male (%)</td>
<td>884 (50.0)</td>
<td>330 (51.4)</td>
<td>318 (50.2)</td>
</tr>
<tr>
<td>Age group (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>16–35 years</td>
<td>885 (50.0)</td>
<td>501 (77.8)</td>
<td>489 (76.8)</td>
</tr>
<tr>
<td>≥36 years</td>
<td>884 (50.0)</td>
<td>37 (5.7)</td>
<td>75 (11.8)</td>
</tr>
<tr>
<td>Ethnicity (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>South Asian</td>
<td>1273 (72.1)</td>
<td>501 (77.8)</td>
<td>489 (76.8)</td>
</tr>
<tr>
<td>Black</td>
<td>140 (8.0)</td>
<td>70 (10.9)</td>
<td>38 (6.0)</td>
</tr>
<tr>
<td>White</td>
<td>259 (14.6)</td>
<td>37 (5.7)</td>
<td>75 (11.8)</td>
</tr>
<tr>
<td>Other</td>
<td>90 (5.1)</td>
<td>36 (5.6)</td>
<td>35 (5.5)</td>
</tr>
<tr>
<td>Origin (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Foreign born (%)</td>
<td>518/876 (59.1)</td>
<td>231/409 (56.5)</td>
<td>107/125 (85.6)</td>
</tr>
<tr>
<td>Index case disease type (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Non-smear positive</td>
<td>1323 (74.8)</td>
<td>501 (77.8)</td>
<td>489 (76.8)</td>
</tr>
<tr>
<td>Pulmonary smear positive</td>
<td>446 (25.2)</td>
<td>70 (10.9)</td>
<td>38 (6.0)</td>
</tr>
<tr>
<td>Household contact (%)</td>
<td>910 (51.4)</td>
<td>367 (56.8)</td>
<td>361 (56.6)</td>
</tr>
<tr>
<td>Partner contact (%)</td>
<td>285 (16.1)</td>
<td>86 (13.3)</td>
<td>153 (24.0)</td>
</tr>
</tbody>
</table>

QFT, QuantiFERON TB Gold-In Tube; TB, tuberculosis.
higher. In conclusion, targeted delivery of 3RH chemoprophylaxis using a QFT based single-step screening strategy is effective and resource efficient for implementation in recent close contacts of active TB below the age of 36 years. Provision of chemoprophylaxis in older contacts of SP TB requires consideration, although effective TB risk stratification may not be achieved with QFT alone for this group.

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Contributors All listed authors meet the criteria for authorship set forth by the International Committee for Medical Journal Editors. PH and GW designed the study, extracted the clinical data and performed the analysis. PM, EWH, MW, JE and GW conceived the screening model and advised on analysis. MB provided microbiological data and wrote the paper with PH and GW. HT leads the contact screening service and resource evaluation of standardized mycobacterial interspersed repetitive-unit-variable-number tandem repeat typing of Mycobacterium tuberculosis. J Clin Microbiol 2008; 46:4986–90.

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