Prospective Comparison of QFT-GIT and T-SPOT.TB Assays for Diagnosis of Active Tuberculosis

Fengjiao Du,1, Li Xie,2, Yonghong Zhang,3, Fei Gao,4, Huibin Zhang,4, Wei Chen,5, Bingqi Sun,5, Wei Sha,6, Yong Fang,6, Hongyan Jia,1, Aiyieng Xing,1, Boping Du,5, Li Zheng,3, Mengqiu Gao,2 & Zongde Zhang1

T-SPOT.TB and QuantiFERON-TB Gold In-Tube (QFT-GIT) tests, as two commercial blood assays for diagnosing active tuberculosis (ATB), are not yet fully validated. Especially, there are no reports on comparing the efficacy between the two tests in the same population in China. A multicenter, prospective comparison study was undertaken at four hospitals specializing in pulmonary diseases. A total of 746 suspected pulmonary TB were enrolled and categorized, including 185 confirmed TB, 298 probable TB and 263 non-TB. Of 32 patients with indeterminate test results (ITRs), age and underlying disease were associated with the rate of ITRs. Furthermore, the rate of ITRs determined by T-SPOT.TB was lower than QFT-GIT (0.4% vs. 4.3%, P < 0.01). When excluding ITRs, the sensitivities of T-SPOT.TB and QFT-GIT were 85.2% and 84.8%, and specificities of 63.4% and 60.5%, respectively in the diagnosis of ATB. The two assays have an overall agreement of 92.3%, but exhibited a poor linear correlation (r^2 = 0.086) between the levels of interferon-γ release detected by the different assays. Although having some heterogeneity in detecting interferon-γ release, both the QFT-GIT and T-SPOT.TB demonstrated high concordance in diagnosing ATB. However, neither of them showed suitability in the definitive diagnosis of the disease.

Tuberculosis (TB) is an airborne-transmitted infectious disease with high morbidity and mortality. In 2015, there were an estimated 10.4 million new TB cases worldwide. China is ranked third, worldwide (after India and Indonesia), with 918 thousand new cases of TB every year.1–3 Early and accurate diagnosis of active TB is critical to the care of TB patients and to control transmission in these high-burden developing countries. Despite incorporation of clinical, radiological, pathological and microbiological examinations, TB diagnosis can still be difficult. Conclusive diagnostic tests, including microbial culture and smear for acid-fast bacilli, are not sensitive enough to identify all the active cases.4

Recently, interferon-γ release assays (IGRAs) have emerged as immunodiagnostic tools to detect tuberculous infection. IGRAs quantify interferon-γ released by T-lymphocytes in response to stimulation by specific antigens encoded in region of difference 1 (RD1) of the Mycobacterium tuberculosis (MTB) genome. Two IGRAs, including an enzyme-linked immunospot (ELISPOT) T-SPOT.TB blood test (T-SPOT.TB; Oxford Immunotec Limited, United Kingdom) and an enzyme-linked immunosorbent assay (ELISA) QuantiFERON-TB Gold In-Tube test (QFT-GIT; Cellestis Limited, Australia), are now commercially available. Several studies have demonstrated that IGRAs may be useful as supplemental tools in the diagnosis of active TB.5–8 Since IGRA methods, interpretation criteria, and study populations varied considerably among published reports, it remains uncertain whether IGRA is suitable for diagnosing active TB or which IGRA is more effective in diagnosing active
TB. Relatively few studies have reported direct comparison between T-SPOT.TB and QFT-GIT assays in diagnosing active TB, most studies being performed with either the T-SPOT.TB or QFT-GIT assay. Furthermore, indeterminate test results (ITRs) of IGRAs could confuse clinical interpretation. If a sample does not respond sufficiently to either specific antigens or the mitogen control, the test result of this sample is deemed as an ITR, yet some evaluations included ITRs in their sensitivity calculations and limited information is available on the risk factors associated with ITRs.

The aim of this study was to assess the diagnostic performance of QFT-GIT and T-SPOT.TB assays in patients with active TB, and compare the concordance between the two diagnostic assays in routine clinical practice in a high TB setting.

Results
Clinical characteristics of participants. In total, 827 participants with clinically suspected pulmonary TB were enrolled in this study. After a follow-up of at least 3 months, 81 patients were excluded from the study, 9 due to lack of data and 72 with no final diagnosis. The remaining 746 participants were ultimately included for T-SPOT.TB and QFT-GIT analyses (Fig. 1). All patients in the study were tested for HIV and all had negative results. Demographic and clinical characteristics of the patients are shown in Table 1. Based on the reference standard, 185 (24.8%) patients were categorized as having confirmed TB, 298 (39.9%) patients were categorized as having probable TB, and 263 (35.2%) patients were categorized as having non-TB. Other organs, in addition to the lung, were affected in 109 (14.6%) of the patients with TB, including lymph node, pleural membrane, bronchus chest wall and bone. Of the 263 patients in the non-TB group, 168 had lung carcinomas, 53 had bacterial pneumonia or a lung abscess, 13 had bronchiectasis and 29 had other lung diseases (Table 1).
Table 2. Univariate and multivariate analysis of risk factors associated with indeterminate IGRA results in 746 patients. IGRAs, interferon-γ release assays. WBC, white blood cells. RBC, red blood cells. ADA, adenosine deaminase. CRP, C-reactive protein. ESR, erythrocyte sedimentation rate. aMedian age 47 years old.

| Category                              | Indeterminate N = 32 (%) | Determinate N = 714 (%) | P-value | OR          | 95% CI       |
|---------------------------------------|--------------------------|-------------------------|---------|-------------|--------------|
| Age ≥ median age³                     | 21 (65.2)                | 303 (42.4)              | 0.010   | 2.841       | 1.339–6.024  |
| Male sex                              | 18 (56.3)                | 448 (62.7)              | 0.458   |             |              |
| Duration of symptom (days)            | 63 (30–170)              | 61 (30–180)             | 0.927   |             |              |
| Underlying disease                    | 18 (56.3)                | 208 (29.1)              | 0.002   | 3.401       | 1.645–6.993  |
| Diabetes mellitus                     | 7 (21.9)                 | 87 (12.2)               | 0.106   |             |              |
| COPD                                  | 4 (12.5)                 | 63 (8.8)                | 0.477   |             |              |
| Connective tissue disease             | 1 (3.1)                  | 9 (1.3)                 | 0.375   |             |              |
| Solid tumor                           | 1 (3.1)                  | 6 (0.8)                 | 0.190   |             |              |
| Virus hepatitis or cirrhosis          | 2 (6.3)                  | 15 (2.1)                | 0.124   |             |              |
| Intestinal obstruction                | 1 (3.1)                  | 6 (0.8)                 | 0.190   |             |              |
| Hyoproteinemia                        | 2 (6.25)                 | 22 (3.1)                | 0.275   |             |              |

Distribution of indeterminate IGRA results and identification of risk factors. Among the 746 TB patients, the T-SPOT.TB assay provided significantly fewer ITRs (0.4% [n = 3]) than did the QFT-GIT assay (4.3% [n = 32], Chi-square value = 24.6, P < 0.001). All 3 patients with ITRs determined by the T-SPOT.TB assay had positive control counts of <20 spot-forming cells (SFCs), while the 32 patients with ITRs determined by the QFT-GIT assay had mitogen minus Nil values of <0.5 IU/mL and antigen minus Nil values of <0.35 IU/mL. For these 32 patients, the T-SPOT.TB assay reported 3 indeterminate, 10 negative and 19 positive results. The 3 patients with ITRs determined by both the QFT-GIT and T-SPOT.TB assays were all greater than 50 years old and two of them had diabetes.

We further compared the basal characteristics and clinical laboratory findings for patients with indeterminate and determinate IGRA results (Table 2). Univariate analysis showed that age (P = 0.010) and underlying disease (P = 0.002) were significantly associated with ITRs. In multivariate analysis, older age [odds ratio (OR) = 2.84, 95% confidence interval (CI) = 1.34–6.02], and underlying disease (OR = 3.40, 95% CI = 1.65–6.99) were independent risk factors for ITRs.

Diagnostic performance of the T-SPOT.TB and QFT-GIT assays for active TB. Of the 743 valid results determined by the T-SPOT.TB assay, the sensitivities were 88.6% (95% CI = 83.3–92.5%) and 83.1% (95% CI = 78.4–87.0%) in the confirmed TB and probable TB groups, respectively, with no significant difference between the two subgroups (P > 0.05). The overall specificity was 63.4% (95% CI = 57.4–69.0%) in non-TB group (Table 3). The PPV, NPV, LR+ and LR− of T-SPOT.TB were 81.1% (95% CI = 74.3–87.8%), 70.3% (95% CI = 63.6–77.1%), 2.33 (95% CI = 1.98–2.74), and 0.233 (95% CI = 0.184–0.294) (Table 4).

Among 714 patients with valid QFT-GIT results, the sensitivities were 88.4% (95% CI = 82.9–92.3%) and 82.5% (95% CI = 77.6–86.5%) in the confirmed TB and probable TB groups, respectively, with no significant difference between the two subgroups (P > 0.05). The overall specificity was 60.3% (95% CI = 54.3–66.3%) in non-TB group (Table 3). The PPV, NPV, LR+ and LR− of QFT-GIT were 79.6% (95% CI = 72.9–86.4%), 68.6% (95% CI = 61.8–75.4%), 2.15 (95% CI = 1.83–2.51) and 0.251 (95% CI = 0.20–0.32) (Table 4).

Both sensitivity and specificity of T-SPOT.TB were slightly higher than those of QFT-GIT, but not reaching statistically significance (all P values > 0.05). We further measured the concordance between the tests using the Kappa index (k value > 0.75, excellent agreement; 0.75 ≥ k ≥ 0.4, fair to good agreement; k < 0.4, poor agreement). In the 714 subjects (excluding the 32 patients with indeterminate results by either of the two assays), both QFT-GIT and T-SPOT.TB were positive in 461 and negative in 197 subjects, and the observed agreement between the tests was 92.3%, with excellent concordance (k = 0.82) (Table 5). When investigating different groups, the agreements were 93.4%, 90.0%, and 93.7% in confirmed TB, probable TB and non-TB groups, respectively.
When combining the T-SPOT.TB and QFT-GIT assays in patients with probable TB, where a positive result was assumed when either test was positive and a negative result was assumed when both tests were negative, the diagnostic sensitivity increased to 87.5% (245/280), the specificity was 58.5% (148/253).

Comparison of interferon-gamma concentration by IGRAs. The numbers of spot forming cells (SFCs) of interferon-γ determined by the T-SPOT.TB assay is shown in Fig. 2. Among 743 patients with valid T-SPOT.TB results, the median numbers of SFCs in the confirmed TB, probable TB and non-TB groups were 89 (Interquartile range (IQR): 22–213), 49 (IQR: 15–164) and 3 (IQR: 0–54) per 2.5 × 10^5 PBMCs, respectively. Both the confirmed TB and probable TB groups had significantly more SFCs than the non-TB group (all P-values < 0.001). The median numbers of SFCs in the confirmed TB group was greater than that in the probable TB group, but the difference did not reach statistical significance (P > 0.05).

### Table 3. The sensitivity and specificity of the T-SPOT.TB and QFT-GIT assays in determining patients with active TB. TB, tuberculosis. *A positive result was assumed when either test was positive and a negative result was assumed when both tests were negative.

| Groups   | Number | T-SPOT.TB | QFT-GIT |
|----------|--------|-----------|---------|
| Confirmed TB | 185 | 88.6 (83.3–92.5) | 88.4 (82.9–92.3) |
| Probable TB | 296 | 83.1 (78.4–87.0) | 82.5 (77.6–86.5) |
| Non-TB | 262 | 63.4 (57.4–69.0) | 60.5 (54.3–66.3) |
| Total | 714 | 85.2 (81.8–88.1) | 84.8 (81.2–87.8) |

### Table 4. Diagnostic performance of the T-SPOT.TB and QFT-GIT assays in active TB. TB, tuberculosis. PPV, positive predictive value; NPV, negative predictive value; LR+, likelihood ratio for positive test; LR−, likelihood ratio for negative value. *There were 3 indeterminate T-SPOT.TB results. **There were 32 indeterminate QFT-GIT results.

| Methods   | Number | Sensitivity % (95% CI) | Specificity % (95% CI) | PPV % (95% CI) | NPV % (95% CI) | LR+ (95% CI) | LR− (95% CI) | Diagnostic odds ratio |
|-----------|--------|------------------------|------------------------|----------------|----------------|--------------|--------------|-----------------------|
| T-SPOT.TB | 743    | 85.2 (82.0–88.3)       | 63.4 (57.4–69.0)       | 81.1 (74.3–87.8) | 70.3 (63.6–77.1) | 2.33         | 0.233       | 9.985                  |
| QFT-GIT   | 714    | 84.8 (81.2–87.8)       | 60.5 (54.3–66.3)       | 79.6 (72.9–86.4) | 68.6 (61.8–75.4) | 2.15         | 0.251       | 8.55                   |

### Table 5. Concordance between the T-SPOT.TB and QFT-GIT assays. *Excluding 32 subjects with indeterminate results by either of the two interferon-γ release assays (T-SPOT.TB and QFT-GIT).

| Groups   | Number | T-SPOT.TB | QFT-GIT | Agreement (95% CI) | OR (95% CI) | Kappa |
|----------|--------|-----------|---------|--------------------|------------|-------|
| Confirmed TB | 181 | Positive | 155 | Negative | 5 | 93.4 (86.6–100) | 62.0 (37.4–221.1) | 0.66 |
| Probable TB | 280 | Positive | 217 | Negative | 14 | 90.0 (83.2–98.8) | 38.8 (17.0–88.2) | 0.65 |
| Non-TB | 253 | Positive | 89 | Negative | 11 | 93.7 (86.9–100) | 239.5 (80.6–711.8) | 0.87 |
| Total | 714* | Positive | 461 | Negative | 26 | 92.3 (85.5–99.1) | 116.4 (67.1–202.0) | 0.82 |

*Excluding 32 subjects with indeterminate results by either of the two interferon-γ release assays (T-SPOT.TB and QFT-GIT).
We further investigated the association between the number of SFCs in the T SPOT.TB and the amounts of released interferon-γ (measured by QFT-GIT). There was an increase of numbers of SFCs when interferon-γ concentrations became higher, with the regression line slope being 7.0972 (P < 0.001). However, regression analysis demonstrated a poor linear correlation (r² = 0.086) between the two tests (Fig. 3).

**Figure 2.** Scatter plots of the SFCs using the T-SPOT.TB assay (A) and the amounts of released interferon-γ using the QFT-GIT (B) assay in the confirmed TB, probable TB and non-TB groups, respectively. The groups were compared using Mann-Whitney tests. SFCs, spot forming cells. PBMCs, peripheral blood mononuclear cells. TB, tuberculosis.

**Figure 3.** Association between the number of SFCs (spot-forming cells) (TB Ag – Nil) in the T-SPOT.TB assay and the amounts of released interferon-γ (TB Ag – Nil) in the QFT-GIT assay among 746 patients. Regression analysis were demonstrated by linear correlation (r²).

**Discussion**

Our findings indicate that, the two IGRAs (T-SPOT.TB and QFT-GIT) with different technical and performance characteristics might have some heterogeneity when used in routine clinical practice, although good agreement in general was observed between the two assays.

To our knowledge, this is the first study to date to directly compare the T-SPOT.TB and QFT-GIT assays for the diagnosis of active TB using the same study participants, in China. In our study, the overall sensitivity of the T-SPOT.TB assay was very close to that of the QFT-GIT assay, within the range from 60% to 89%7,8,16–19. And parallel with a recent meta-analysis, was 84.0% (95% CI 81.4%–86.4%) for T-SPOT.TB and 84.2% (95% CI 81.1%–87.0%) for QFT-GIT20. However, the sensitivity of T-SPOT.TB (85.2%) in our study was lower than the few studies performed in China by Zhang et al. (94.7%)7, Feng et al. (94.7%)21, and Kang et al. (93.2%)22. From the ten reported studies evaluating T-SPOT.TB in China, the combined sensitivity was 88%23. The sensitivity of QFT-GIT
(84.8%) in our study was higher than the reported range in the earlier studies from highly endemic countries, Xia
et al. (80.9%) in China8, 81% in India24, and 64.0% in Gambia25.

This multicenter, prospective study was conducted at four hospitals specializing in pulmonary diseases, with
high experienced technics and the sophisticated laboratory conditions. We simultaneously compared T-SPOT.TB
and QFT-GIT on the same blood samples from the same participants for each test. This may be caused the high
concordance of sensitivity between the two IGRAs. Furthermore, The QFT-GIT is based on the ELISA method,
while T-SPOTT.B assay is based on the ELISPOT method. The sensitivity of QFT-GIT is strongly affected by
immunosuppressed status with only a slight effect of antigenic load, whereas the sensitivity of T-SPOTT.B is
strongly affected by the antigenic load with only a slight effect of immunosuppressed status11,26. The present study
enrolled specified patients with active TB with nonimmunosuppressed conditions, seems to reduce the differences
between the blood tests might provide slightly different results in generalization. In addition, both T-SPOTT.B
and QFT-GIT had imperfect sensitivities and poor negative predictive values (70.3% and 68.6) could give rise to
potential problems when using negative results as a “rule-out” criterion in the diagnosis of active TB. Previous
studies have interpreted multiple factors that may affect IGRAs15,27,28. Careful interpretation of negative IGRA
results is necessary for patients who are older, over-weight, have HIV coinfection or are receiving TB treatment.
Caution is also needed in interpretation of results based on the patient's immunosuppression status and certain
HLA-genotypes.

The diagnosis of TB in patients with negative bacteriological results remains a problem in clinical settings.
In our subgroups, the sensitivities of both tests in confirmed TB and probable TB groups were not significantly
different. When combining the two tests, the overall sensitivity rose to 87.5% with minimal loss of specificity
(from 60.5 to 58.5%) in the probable TB group. Thus, the commercially available versions of the T-SPOTT.B
and QFT-GIT assays may be of complementary diagnostic value for the assessment of bacteriologically negative
TB3,29.

The low specificities of the two assays, 63.4% for T-SPOTT.B and 60.5% for QFT-GIT, limit their usefulness
in routine clinical practice, at least in a setting highly endemic for TB such as in China20,30. As only a small
minority of lymphocytes in the human body circulate in the blood, neither immunological test could accurately
distinguish between untreated active TB and latent tuberculosis infection (LTBI) when IGRAs were performed
on the blood cells31. Latent TB has a high prevalence, about 19% in rural China based on the QFT-GIT assay
and likely contributes a decrease in the specificity of the IGRAs32. In this study, specificities of the tests were poor
for active TB compared with previously reported data34. This study was designed to evaluate the diagnostic validity
of T-SPOT.TB and QFT-GIT in routine clinical practice, and thus focused on unselected patients with suspected
active TB. Therefore, in this setting, the diagnostic validity tends to be lower than that in studies in which healthy
people are enrolled as negative controls and patients with active TB as positive controls. Furthermore, although
the IGRAs have been confirmed to be more sensitive and specific than the tuberculosis skin test (TST) since the
assays use M. tuberculosis specific antigens, we should not ignore the fact that patients infected with M. kansasii,
M. szulgator M. marinum may also lead to positive results in IGRAs20,28,33.

One disadvantage of interferon-γ assays is that they can yield ITRs. Previous studies found that ITRs were
common (ranging from 0% to 20%) when using QFT-GIT assay14,15,29,30,34. However, the majority of them
were associated with immunosuppression or old age. In this study, the rate of ITRs was significantly higher for
QFT-GIT than that for T-SPOTT.B. Of the 32 patients with indeterminate QFT-GIT results, only 3 showed
indeterminate results by T-SPOT.TB test. Furthermore, most indeterminate results of both tests were caused by insuf-
ficient response to the positive control antigens. It can be speculated that the patients with weak immune system
response to the positive control yield a response that is insufficiently strong to be measured11,13. The main differences
between the QFT-GIT and T-SPOTT.B lie in the specimens used (whole blood versus mononuclear cells).
The use of whole blood by the QFT-GIT may lead to the presence of insufficient mononuclear cells producing
various kinds of cytokines, so that indeterminate QFT-GIT result was more common than for T-SPOTT.B. As for
clinical characteristics, patients showing indeterminate results on both tests were most frequently elderly patients
or patients with severe underlying diseases such as diabetes mellitus. In our study, the number of indeterminate
results was small and there were no patients who were serum positive for HIV. As there are no reports demonstrat-
ing a decrease in the functional activity of mononuclear cells for these patients suspected of TB disease, it will
be necessary to investigate the functional activity of mononuclear cells of these patients in future20,28,34.

Although both the T-SPOT.TB and QFT-GIT assays measure T-cell interferon-γ responses to similar M.
tuberculosis-specific antigens over a 16- to 24-h incubation period, they are based on different technology plat-
forms. The mycobacterial antigen load (which reflects bacterial load) is positively correlated with effector T cell
levels13,35. Similar to previous studies, in the present study, the level of interferon-γ release (SFCs by T-SPOTT.B
measured in the QFT-GIT by the regression analysis, we obtained a
poor correlation (r² = 0.086) between the two assays (Fig. 3). A few factors should be considered to interpret this
poor correlation. T-SPOTT.B requires a specific number of peripheral blood mononuclear cells and measures the
number of interferon-γ-secreting T cells via ELISPOT assay, whereas QFT-GIT measures the concentration of interferon-γ via an ELISA using specific volume of whole blood. This indicate that we cannot rule out the possibility that a small number of T cells can produce a great deal of interferon-γ in some patients. ELISA (measured as "IU/mL") is more vulnerable to the influence of incubation time than ELISPOT (measured as "SFCs/2.5 x 105
PBMCs"). In addition, the T-SPOTT.B method use of separate mixtures of ESAT-6 and CFP-10 synthetic pep-
tides as M. tuberculosis-specific antigens, compared to the QFT-GIT use of a single mixture of synthetic ESAT-6,
CFP-10, and TB7.7 peptides could also lead to the poor correlation. Previous studies have reported that the use of both techniques simultaneously can contribute to improving the knowledge of TB immunity. Therefore, the results of poor correlation require further investigation to elucidate possible clinical implications.

To our best knowledge, this is the largest study to date to directly compare the two IGRA.s currently commercially available for the diagnosis of active TB. However, we did not enroll LTBI subjects in the control group, which might have some impact on specificity, since the two IGRA.s cannot distinguish between LTBI and active TB. In addition, immunodeficient patients should be included to finalize the study results.

Furthermore, QuantiFERON-TB Plus (QFT-Plus) as an updated version of QFT-GIT has been on the market in Europe and the United States, but not commercially available in China. QFT-Plus includes two tubes, TB1 and TB2 with MTB antigens to elicit a specific immune response. TB1 is designed to induce a specific CD4 T-cell response. TB2 contains newly designed peptides stimulating interferon-γ production by both CD4 and CD8 T cells. The additional peptides for eliciting CD8 T-cell responses may reduce the indeterminate results of diagnosing active TB.

In conclusion, while there were some differences in the performance of the T-SPOT.TB and QFT-GIT assays, the two commercialized IGRA.s have similar sensitivities to aid in the diagnosis of active TB. When combined, the two assays may be of complementary diagnostic value for probable TB. However, the high false positive rates of these tests limit their usefulness in routine clinical practice in China, where the prevalence of LTBI is high. Therefore, these assays should not be used alone to rule out or rule in active TB cases, and further modification is needed to improve their accuracy.

Methods
Patients and setting. This multicenter, prospective comparison study was conducted at Beijing Chest Hospital, Shanghai Pulmonary Hospital, Inner Mongolia Autonomous Region Fourth Hospital and Shenyang Chest Hospital in China. The four hospitals are situated in the south, north, east, and center of China and specialize in the diagnosis and treatment of pulmonary diseases, especially TB and lung cancer. Between June 2012 and November 2013, a total of 830 patients with presumed pulmonary TB who underwent valid T-SPOT.TB and QFT-GIT tests before the onset of anti-TB therapy were enrolled in the study. Patients gave written informed consent and were included if they had any clinical symptoms, signs or radiographic evidence of active TB. Patients were excluded if they had a history of previous TB or TB contact or had received anti-TB therapy before enrollment.

Medical records were collected on age, gender, underlying disease, HIV serology and duration of illness before hospitalization. The ‘duration of illness before hospitalization’ was defined as the length of time from the onset of TB clinical symptoms to the time of admission to hospital. Routine clinical, microbiologic, histopathological, radiological examinations were also performed and collected over a follow-up period of at least 3 months. The microbiologic examinations included, at a minimum, three sputum smears and one sputum culture. All patients were interviewed and a questionnaire was completed to obtain epidemiological data.

This study was performed in accordance with the guidelines of the Helsinki Declaration and was approved by the Ethics Committee of the Beijing Chest Hospital, Capital Medical University.

Definitions and diagnosis. All participants were classified into one of 3 groups depending on their clinical manifestations, bacteriological, biochemical examinations, histopathological examination and responses to anti-TB therapies. The first group, designated as confirmed TB, consisted of patients with microbiologically confirmed pulmonary TB who tested positive three 3 times in sputum smears or culture analyses. The second group was designated as probable TB, consisting of patients who were classified as having pulmonary TB based on clinical and radiological findings, but lacking microbiological evidence of M. tuberculosis infection. All patients with confirmed TB or probable TB responded clinically to antituberculous chemotherapy during follow-up and were categorized as having active TB. The final group, designated as the control group (non-TB), contains patients categorized as not having active TB if other diagnoses were made or if clinical improvement occurred without recent anti-TB therapy.

T-SPOT.TB assay. The T-SPOT.TB (Oxford Immunotec Limited, UK) was performed according to the manufacturer’s instructions in Class II Biological Safety Cabinet (ESCO A2, Singapore). Six mL of heparinized peripheral blood samples were collected and processed within 2h of collection. Peripheral blood mononuclear cells (PBMCs) were isolated and adjusted to a concentration of 2.5 × 10^6/mL and then the wells were stimulated with 50μL of each of phytohemagglutinin (positive control), secretory antigenic target (ESAT)-6, culture filtrate protein (CFP)-10, and AIM® V medium (Invitrogen, USA) (negative control). The procedure was performed in plates pre-coated with anti-interferon-γ antibodies at 37 °C for 16 to 20 hours. After application of the alkaline phosphatase-conjugated second antibody and chromogenic substrate, the number of spot forming cells (SFCs) in each well was automatically counted with a CTL ELISPOT system (CTL-ImmunoSpot® S5 Versa Analyzer, USA). The criteria for positive, negative, and indeterminate outcomes were recommended by the manufacturer.

QuantiFERON-TB Gold In-Tube assay. The QFT-GIT (Cellestis Limited, Australia) was performed according to the manufacturer’s instructions in Class II Biological Safety Cabinet (ESCO A2, Singapore). Three mL peripheral blood was drawn from each patient on the day of enrollment. The blood was collected in 3 special tubes: 1 coated with M. tuberculosis-specific peptides (TBAg: ESAT-6, CFP-10, and TB 7.7), 1 coated with mitogen as a positive control, and 1 without antigen coating as a negative control (Nil). Within 8h of blood sampling, the tubes were incubated for 16–24 h at 37 °C, centrifuged, and stored at −20 °C until assayed (within 7 days). The plasma interferon-γ concentration was measured by ELISA. The test results were determined as negative, indeterminate, or positive (cut off at 0.35 IU/mL) according to the manufacturer’s software.
**Statistical analysis.** Data analysis was performed using SPSS, Version 17.0 (SPSS, Inc, Chicago, IL, USA). Categorical data were compared by Pearson’s Chi-square or Fisher’s exact test. Odds ratio (OR) and multiple logistic regression analysis were used to calculate crude and adjusted risk, and model building was performed backward using the chance criteria for variable selection. Covariates that were considered biologically important were forced into the model irrespective of statistical significance. In addition, sensitivity, specificity, positive predictive value (PPV), negative predictive value (NPV), likelihood ratio positive (LR+), and likelihood ratio negative (LR−) were calculated to evaluate diagnostic performance for the T-SPOT.TB and QFT-GIT assays. Ninety five percent confidence intervals (95% CI) were estimated according to the binomial distribution. Continuous variables were compared using nonparametric Mann-Whitney U test. All P values reported were calculated two-tailed with statistical significance set to P < 0.05.

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**Author Contributions**

Zongde Zhang, Mengqiu Gao, and Li Zheng were the principal investigators who conceived the study and obtained financial supports. Zongde Zhang, Mengqiu Gao, Fengjiao Du, Li Xie, Fei Gao, Huibin Zhang, Wei Chen, Bingqi Sun, Wei Sha and Yong Fang designed the study. Li Xie, Huibin Zhang, Wei Chen, and Yong Fang were responsible for the recruitment of participants, collection of clinical information, and sample preparation. Fengjiao Du, Fei Gao, Bingqi Sun, Wei Sha, Hongyan Jia, Aiying Xing and Boping Du performed the T-SPOT.TB and QFT-GIT assays. Fengjiao Du, Yonghong Zhang and Li Zheng conducted data management and performed the statistical analyses. Fengjiao Du and Li Xie interpreted the results and drafted the manuscript. Zongde Zhang, Mengqiu Gao, Yonghong Zhang, Fengjiao Du, Li Xie, Li Zheng, Fei Gao, Huibin Zhang, Wei Chen, Bingqi Sun, Wei Sha, Yong Fang, Hongyan Jia, Aiying Xing and Boping Du approved the final version of the manuscript.

**Additional Information**

**Competing Interests:** The authors declare no competing interests.

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