

First evaluation of QuantiFERON-TB Gold Plus performance in contact screening

L. Barcellini ^{1*}, E. Borroni¹, J. Brown², E. Brunetti³, D. Campisi⁴, P. Castellotti⁴, L.R. Codecasa⁴, F. Cugnata⁵, C. Di Serio⁵, M.Ferrarese⁴, D. Goletti⁶, M. Lipman², P.M.V. Rancoita⁵, G. Russo¹ M. Tadolini⁷, E.Vanino⁷ and D.M. Cirillo^{1§}

¹ Emerging Bacterial Pathogens Unit, Division of Immunology and Infectious Diseases IRCCS San Raffaele Scientific Institute, Milano Italy

² Royal Free London NHS Trust and UCL Respiratory, Division of Medicine, University College London, UK.

³ Department of Infectious Diseases, San Matteo Hospital Foundation- University of Pavia, Pavia, Italy

⁴Regional TB Reference Centre and laboratory, Villa Marelli Institute/Niguarda Ca' Granda Hospital, Milan, Italy

⁵ University Centre of Statistics for Biomedical Sciences (CUSSB), Vita-Salute San Raffaele University, Milano Italy

⁶ Translational Research Unit, Epidemiology Department, National Institute for Infectious Diseases, Rome, Italy

⁷ Infectious Diseases Unit, Department of Medical and Surgical Sciences, Alma Mater Studiorum University of Bologna, Italy

[§]Corresponding Author:

Daniela M. Cirillo Emerging Bacterial Pathogens Unit San Raffaele Scientific Institute Via Olgettina 58, 20132 Milano, Italy Phone ⁺39 02 2643 7947 <u>cirillo.daniela@hsr.it</u> Support: QIAGEN (Hilden, Germany) provided the QuantiFERON-Pus Kits free of charge;

Take home message: QFT-Plus improved the diagnostic accuracy for LTBI in the setting of contact screening

Total word count: 3280

- 1 ABSTRACT
- 2
- 3

4

5 evaluated for the first time the performance of a new type of interferon- γ release assays, 6 QuantiFERON-TB Plus that includes an additional antigen tube (TB2) stimulating both CD4⁺ 7 and CD8⁺ T-cells in contacts of tuberculosis patients. 8 9 Materials and Methods: Contacts were screened for latent tuberculosis infection by 10 tuberculin-skin-test, QuantiFERON-TB Plus (QFT-Plus) and QuantiFERON-TB Gold in 11 Tube (QFT-GIT). 12 13 Results: In 119 TB contacts, the overall agreement between QFT-Plus and QFT-GIT was 14 high, with a Cohen's kappa of 0.8. Discordant results were found in 12 subjects with negative 15 QFT-GIT and positive QFT-Plus results. In analyses of markers of tuberculosis exposure and 16 tests results, the average time spent with the index case was the strongest risk factor for both 17 tests' positivity. The difference in interferon- γ production between the two antigen tubes 18 (TB2-TB1) was used as an estimate of CD8⁺ stimulation provided by the TB2. TB2-TB1 19 values >0.6ml/IU were significantly associated with proximity to the index case and European 20 origin. 21 22 Conclusion: QuantiFERON-Plus has a stronger association with surrogate measures of TB 23 exposure than QFT-GIT in adults screened for LTBI. Interferon-y response in the new antigen 24 tube used an indirect estimate of specific CD8⁺ response correlates with increased M. 25 tuberculosis exposure suggesting a possible role in identifying individuals with recent 26 infection. 27 28 29 30 **Total words count: 200** 31 32 33 **Key words:** latent tuberculosis infection, Interferon-γ release assay, CD8⁺ T-cell, 34

Rationale: Identifying latently infected individuals is crucial for tuberculosis elimination. We

35 36

37 INTRODUCTION

Despite the progress recently made in tuberculosis (TB) control at a global level, the decline in TB incidence is much slower than that needed to achieve TB elimination by 2050[1]. Identifying and treating symptom-free people who are truly latently infected with *M.tuberculosis* (Mtb) is key to achieving this[2][3]. The current global burden of latent infection is uncertain, although it has been suggested that one third of the world's population may be latently infected with Mtb[4].

44 Although they show no sign of disease, individuals with latent tuberculosis infection (LTBI) 45 are at risk of reactivating and up to 10% of them may develop active disease in their lifetime[5]. 46 This risk is highest in the first 2 years following infection. Preventive treatment of recently-47 infected individuals reduces this[6]. However isoniazid preventive therapy is not optimal for a 48 large-scale implementation program, and the current LTBI diagnostic tests - Tuberculin Skin 49 Test (TST) and Interferon- γ release assay (IGRAs)- have significant limitations. TST may be 50 falsely positive due to sensitization by environmental mycobacteria and BCG-vaccination[7]. 51 In recent decades IGRAs measuring the INF-y concentration after in vitro whole blood 52 stimulation with peptides from the RD-1 region of the Mtb genome were developed to improve 53 specificity of the diagnosis[8]. IGRAs are a useful indicator of Mtb exposure as their specificity 54 is very high (97%)[9]. However, like TST, they lose sensitivity in immune-compromised 55 individuals and children[10][11]; they identify both recent and past infection and they are poor 56 at predicting LTBI subjects who are at greater risk of developing disease (positive predictive 57 value IGRA 2,7% TST 1,5%)[12]. As a result when currently available diagnostic tests are used 58 to guide the administration of preventive therapy, the number needed to treat to prevent one 59 case of TB is too high to allow a large-scale preventive program. Different approaches have 60 been described in the literature to help discriminate those at greater risk of active TB 61 development. The use of $INF-\gamma$ response to the latency antigen Heparin-Binding-62 Haemoagglutinin (HBHA)[13][14], immunoprofiling[15][16], gene expression pattern (i.e. IL-63 13 and AIRE)[17][18] and proportion of peripheral blood monocytes[19] have been studied as 64 possible biomarkers for incipient TB. However all of these approaches are still confined to 65 research fields and currently have minimal impact on patient management.

66 QuantiFERON-TB Plus (QFT-Plus) is a new generation of QTF-Gold In Tube (QFT-GIT) [5]

67 that includes an additional antigen tube (TB2). The TB1 tube contains ESAT-6- and CFP-10-

derived peptides (TB-7.7, present in QFT-GIT, has been removed), designed to elicit cell-

69 mediated immune responses from CD4⁺ T-helper lymphocytes. TB2 contains new peptides

able to stimulate IFN- γ production by both CD4⁺ and CD8⁺ T-cells [20].

71 Evidence supports the important contribution of CD8⁺ T-cells in host defense against Mtb by 72 both cytokine secretion and cytotoxic activity[21]. Firstly a positive correlation between 73 specific $CD8^+$ T cells and increased mycobacterial load has been found in peripheral blood ex 74 vivo[22]. Day et al. reported that more than 60% of individuals with smear-positive TB had 75 detectable CD8⁺ T cells response compared with 38% and 20% of smear-negative and LTBI 76 respectively. Consistent with this paradigm a higher prevalence of Mtb-specific CD8⁺ T cells 77 have been reported in smear-positive versus smear-negative patients and in PTB compared with 78 EPTB[23]. In addition, a positive correlation between the $CD8^+$ T cells response against TB 79 antigens and a recent exposure to Mtb have been found. Recent contacts of active TB patients, 80 independent of their response to QTF, have a greater CD8⁺ T cell response compared to other 81 study groups (active TB patients, health care workers, BCG-vaccinated healthy controls)[24]. 82 This is in agreement with findings observed in a cattle model where a CD8⁺ T cell response is 83 present at the onset of infection.[25]

The INF-γ release assays currently in use primarily elicit a CD4⁺ response, but emerging data
provide a good rationale for also measuring specific CD8⁺ T cell responses and in particular
to further investigate the association between CD8⁺ T cells and risk of disease progression.

In the present study we evaluate the performance characteristics of the new QFT-Plus assay in TST-positive contacts with recent exposure to people with confirmed active tuberculosis, assessing the use of QFT-Plus head-to-head with the previous QFT-GIT. In addition, we investigate for the first time the significance and the possible use of the CD8⁺ INF- γ response provided by the second newly-added antigen tube.

92

93 MATERIAL AND METHOD

94 Study setting and participants

We conducted a cross-sectional study at Villa Marelli-Niguarda Hospital. TB incidence in
Milan is of 16.6 new cases per 100.000 persons year (in 2011)[26], three-times higher than the
Italian national average. From November 2014 to June 2015 we prospectively recruited TSTpositive (TST≥5mm) contacts of notified active TB cases sent by the local public health
services to be screened for LTBI.

100 Contacts were excluded if aged less than 18 years old, a previous positive TST was documented,

101 preventive TB treatment was prescribed or past TB history was reported. Informed written

- 102 consent was obtained from each study subject.
- 103 Contacts reporting mild or severe immunosuppression (diabetes mellitus, chronic kidney

- 104 disease, HIV, malignancy, immunosuppressive medications) were included.
- 105 The study was approved by the Ethics committee^{*}.

106 Contact screening strategy was based on the National Institute of Clinical Excellence TB 107 guidelines 2011 [27] and Italian guidelines which recommends retesting those with positive 108 TST results using an IGRA as confirmatory test. At the contact's first visit health status was 109 established by clinical examination and chest X-ray. Further information on the country of birth, 110 immigration status, nature of the contact to the source case, BCG-vaccination status (if details 111 were unclear inspection of BCG-vaccination scar was performed by trained healthcare-112 assistants), and clinical history were obtained through personal interviews. When clinical 113 suspicion persisted, chest CT scan and sputum sample analyses were requested.

All patients also underwent testing in line with recommended routine screening as part of contact investigation. Thus, TST-positive contacts who tested negative to a first QFT-GIT analysis were retested with QFT-GIT after 10-12 weeks to exclude delayed conversion.

117 Blood samples were obtained for QFT-GIT, QFT-Plus and HIV testing from all subjects 118 providing informed consent. QFT-GIT currently in use in clinical practice was performed at 119 Niguarda Microbiology service while QFT-Plus was carried out in the Emerging Bacterial 120 Pathogen Laboratory at San Raffaele Hospital. The QFT-Plus and QFT-GIT tests were 121 performed according to the manufacturer's instructions. Peripheral blood samples for the two 122 tests were obtained simultaneously directly into the QFT tubes and processed within 4 h. Test 123 interpretation for both QFT-Plus and QFT-GIT was performed according to the manufacturer's 124 instruction manual. QFT-GIT results were recorded positive if the antigen response were >0.35125 ml/UI above the negative control response. Positivity (antigen response >0.35 ml/UI above the 126 negative control response) of a single antigen tube (either TB1 or TB2) was sufficient to score 127 the QFT-Plus test as positive.

128

129 Ascertainment of exposure

We assessed different factors as surrogate markers of Mtb exposure. The aggregate exposure time of contacts prior to the diagnosis of their respective source case was established by recording the extent of the contact during a typical week. TB contacts were categorized according to proximity to the index case[28]: we considered them to be "high proximity" if contacts and case patient were sharing routinely the same bedroom and lower proximity if contacts and case patient were sleeping in a different bedroom in the same house or in a different

^{* (}GO/URC/ER/mm prot. n.82/DG, 26 Feb 2010 and successive amendments)

house. Sputum smear positivity of the index case was also assessed as TB case related riskfactor.

138 Statistical Analysis

139 The agreement between QFT-GIT and QFT-Plus was evaluated by computing the overall 140 percent of concordant results and Cohen's kappa coefficient with 95% Confidence Interval (CI). 141 Univariate logistic regression and backward stepwise multivariate logistic regression models 142 were used to identify factors associated with positive test results. The variables considered in 143 the analyses were: gender, whether the country of birth was an endemic area of TB and whether 144 it was European, BCG vaccination, immunocompromised status, smear status of index case, 145 average time spent per week with the index case and place of sleeping with respect to the index 146 case. The same analysis was performed for the variable denoting whether the differences 147 between QFT-Plus TB2 and QFT-Plus TB1 was greater than the cut-off 0.6 IU/ml (as described 148 in the Results). The level of significance considered was 5%. All statistical analyses were done 149 using R statistical software (version 3.2.3).

150

151 **RESULTS**

A total of 119 Mtb-exposed individuals with positive TST (5mm) were investigated. Of these, 39 were contacts of a smear-negative culture-positive TB case, and 69 of a smear-positive culture positive index case. Participants had a median age of 38 years (25-75 percentile: 30-79), more than half (n=61, 51.26%) were non-European-born, 82 (78.85%) were BCGvaccinated and 11 (9.24%) were immunocompromised subjects. Demographic characteristics of the cohort are shown in the Table 1.

158 Agreement between QFT-Plus and QFT-GIT

159 Sixty-eight out of 119 (57.1%) contacts were QFT-Plus positive. 64 subjects were positive in 160 both antigen tubes, 2 were positive to TB1 only and 2 were positive to TB2 only. Fifty-six of 161 119 TST-positive contacts were positive to QFT-GIT. The overall agreement between the two 162 IGRAs was high, with a Cohen's kappa of 0.8 (95% CI 0.69-0.91). The two tests gave 163 concordant results for 107 (89.9%) subjects (see Table 2). Discordant results were found in 12 164 subjects: they all scored negative to the QFT-GIT and positive to the QFT-Plus. Discordant 165 results between the two IGRAs included the 4 contacts with a single tube QFT-Plus positivity. 166 Moreover, contacts with IGRAs discordant results had overall low INF-y responses but not as

- 167 low to be considered borderline results (median: TB1-Nil=0.83 IU/ml, TB2-Nil=0.73 IU/ml).
- 168 The characteristics of subjects with discordant results are shown in Table 3. Only one of the 12
- 169 contacts with QFT-Plus positive and QFT-GIT negative results had a TST response less than
- 170 10mm (7mm). Globally, the median TB1 QFT-Plus antigen IFN-γ level (TB1-Nil) was 0.74
- 171 IU/ml, whereas the median TB2 QFT-Plus antigen IFN-γ level (TB2-Nil) was 0.67 IU/ml, as
- 172 reported in Table 2.

173 As per the Italian guidelines, contacts of TB cases with initial positive TST results who tested 174 negative to a first QFT-GIT analysis, were re-tested with QFT-GIT at 10-12 weeks. At the post-175 exposure follow-up, two contacts converted to QFT-GIT positive results. Both of them were part of the 12 contacts who initially showed QFT-plus-positive/QFT-GIT-negative discordant 176 177 results (Table 2). One of them had a strong QFT-GIT positivity (>10 ml/IU) at 10 weeks post-178 exposure follow-up; while the second case reported a QFT-GIT of 0.5 ml/UI after 6 month of 179 isoniazid preventive therapy (decision to treat was based on the strong TST positivity and the 180 proximity of contact with the index case). In both cases the TB2 INF- γ response was greater 181 than that found in TB1.

182 Independent predictors of QFT-Plus and QFT-GIT positivity

183 For both QFT-GIT and QFT Plus test, the univariate odds ratios of being positive for different 184 possible surrogate markers of increasing exposure to Mtb is presented in Table 4. Contacts 185 reporting that they had spent more than 12 hours per day with the index case were significantly 186 more likely to be both QFT-GIT and QFT-Plus positive, compared to contacts spending 1-4 187 hours per day with the index case. For a subject with an exposure time > 12 hours, the odds of 188 a positive test were 6 times higher by QFT-GIT and 14 times higher by QFT-Plus. Both test 189 results were significantly more likely to be positive in subjects with closer sleeping proximity 190 to the patient (same house versus different house). The odds of being QFT-GIT positive for 191 subjects sleeping in the same house of the index case were approximately 4 times (different 192 rooms: 3.79; same room: 3.98) higher than for those sleeping in a different house, whereas their 193 odds of being QFT-Plus positive were approximately 6 times (different rooms: 5.78; same 194 room: 5.65). The results of the backward stepwise multivariate logistic regression analysis are 195 presented in Table 5. Only the variable indicating whether a contact spent on average more than 196 12h per day with the index case remained significantly associated with a positive QFT-GIT 197 result (OR: 4.63; 95% CI: 2.05-10.47) and a positive QFT-Plus result (OR: 6.98; 95% CI: 2.86-198 17.02).

Predictors for CD8 T-cell stimulation

200 To assess the specific contribution of CD8⁺ T cells, we subtracted the quantitative value of the 201 first antigen tube expressed in IU/ml (TB1), which stimulates the CD4⁺ population only, from 202 the value provided by the second antigen tube (TB2), in which a combined CD4⁺ and CD8⁺ T 203 cell stimulation occurred. We used a difference of 0.6 IU/ml to define positive results in order 204 to reduce the bias of the intrinsic variability of the test[29]. Eighteen contacts out of 119 205 (15.13%) had a difference between TB2 and TB1 greater than 0.6 IU/ml. Univariate logistic 206 regression was used to identify factors associated with differences between TB2 and TB1 > 0.6207 IU/ml (Table 4). This method identified sleeping in the same room compared to sleeping in different houses (OR: 4.34; 95% CI: 1.37-13.81), and European origin (OR: 3.24; 95% CI: 1.07-208 209 9.75) to be to be significantly positively associated with a greater TB2 response. These 210 associations persisted in the multivariate analysis, shown in Table 6.

211

212 **DISCUSSION**

We provide the first evaluation of QFT-Plus assay alongside the previous version QFT-GIT ina cohort of TST-positive contacts of active TB cases.

- 215 Positive results from QFT-Plus were associated with surrogate markers of increasing recent 216 exposure to Mtb. Paired comparison between QFT-GIT and QFT-Plus shows an overall good, 217 but not complete agreement. Furthermore the overall INF-y response in QFT-Plus 218 positive/QFT-GIT negative contacts was in the majority of cases out of the uncertainty zone 219 for test interpretation [29], suggesting that differences between the tests are not due to test 220 variability. Of note, the disagreement between the two tests all goes in the same direction, with 221 a total of 12 TST-positive contacts positive with the new QFT-Plus and negative to QFT-GIT. 222 With no gold standard for LTBI to refer to, it is difficult to assess whether the discordant results 223 found during the contact screening are attributable to the higher sensitivity of the QFT-Plus 224 test. If the TST were taken as the reference test for LTBI, this would mean that the proportion 225 of TST-positive contacts confirmed by the IGRA test is increased by 17% when using the QFT-226 Plus compared to QFT-GIT. False positivity with TST is mainly due to sensitization by BCG-227 vaccination[7]. QFT-Plus specificity in a BCG-vaccinated population has not been investigated 228 yet, however we found that QFT-Plus is not associated with BCG-vaccination both in univariate 229 and multivariate analysis. Moreover only one of the 12 contacts with QFT-Plus positive and 230 QFT-GIT negative result had a TST response less than 10mm while another showed an intense 231 TST positivity which is less likely to be the result of previous vaccination[7]
- 232 Recent findings suggest that the discordance between IGRAs and TST in recently-exposed

individuals may be related to delayed conversion of IGRAs relative to TST[30][31]. In this
study we find that most of the discordant cases (QFT-GIT negative/QFT-Plus positive) show
intense TST positivity; moreover, we reported a shorter period of conversion for QFT-Plus
compared to QFT-GIT at least in two individuals of our cohort. These results suggest that QFTPlus may be more sensitive in detecting new or recent infection with Mtb than the QFT-GIT.

Our data demonstrate that risk factors for test positivity were the same for both IGRAs. QFT-Plus showed stronger associations with surrogate measure of recent exposure than QFT-GIT both in univariate and multivariate analysis The average time spent per day with the index case had the strongest association with test positivity.

242 We investigated for the first time the difference in INF-y production between the two QFT-Plus 243 tubes and surrogate markers of increasing exposure. TB2-TB1 differential values were used as 244 an indirect estimate of specific CD8⁺ stimulation with the newly added antigens. A cut-off value 245 was set at 0.6 ml /IU in order to exclude small variations due to inter-test variability[29]. Positive TB2-TB1 differences (>0.6ml/IU) were significantly associated with sleeping 246 247 proximity to the index case with an odds ratio comparable to the one obtained in the analysis 248 of QFT-GIT and QFT-Plus (sleeping in the same room compared to sleeping in different houses 249 OR: 4.34; 95%CI: 1.37-13.81). Moreover, European origin (OR: 3.24; 95%CI: 1.07-9.75) was 250 significantly associated with TB2-TB1 > 0.6ml/IU, while it was not statistically significant for 251 the QFT-GIT and the QFT-Plus results.

As individuals from European countries have a low risk for Mtb exposure, these findings are consistent with the hypothesis that the difference in response between the TB1 and TB2 tubes could be used as a surrogate marker of recent exposure (linked to the specific index case exposure), and not to previous cumulative Mtb exposure. A recent flow-cytometry study reported a positive correlation between the CD8⁺ T cells response against the QFT-GIT antigens and recent exposure to Mtb in contacts of active TB patients compared to controls (active TB patients, health care workers, BCG-vaccinated healthy controls)[24].

Tests currently used for Mtb infection diagnosis do not reflect CD8+ T cell cytokine production[32], however results reported in previous flow-cytometry studies and our own findings provide a strong rationale for measurement of Mtb-specific CD8+ T cell response. If validated, this may prove to be a surrogate marker of recent infection which, having the highest risk of progression to active TB, may enable QFT-Plus to distinguish recent infection from long lasting reactivity and hence allow better targeted delivery of preventive therapy.

Mtb-specific CD8⁺ T cell have been more frequently detected in individuals with active TB when compared with LTBI and correlated with increasing antigenic burden[21][23][22][33]

267 [34], suggesting that the presence of CD8⁺ T cells in a small proportion of latently infected 268 individuals may be predictive of Mtb active replication and more likely disease progression[22]. 269 Consistent with these results, in a previous study we found that the difference in responses 270 between the QFT-Plus tubes may positively correlate with increasing antigenic load in active 271 TB patients, as it was significantly more common in smear-positive versus smear-negative 272 active TB patients[35]. In the present study, we observed a greater TB2 antigen response (TB2-273 TB1 difference >0.6ml/UI) in 18 (15.13%) individuals, all QFT-Plus positive. We speculate 274 that the small subgroup of latently infected contacts with TB2-TB1 difference >0.6ml/UI have 275 higher antigenic burden. However, to date, we do not have the tools to directly assess Mtb 276 antigenic burden, as current LTBI tests rely on the (indirect) measurement of a specific immune 277 response.

278 Our study has limitations. The foremost of these was the sample size, which comprises 119 279 subjects. Moreover because of the lack of gold standard tests for LTBI, we were unable to 280 adequately resolve the discordance between QFT-GIT and QFT-Plus. In addition, TST-281 negative contacts were not recruited in our sample and a full evaluation of the test would benefit 282 of their presence. Finally, the positive predictive value of the test and of the new parameter, the 283 difference between the two antigen tubes, needs to be properly assessed in a longitudinal cohort. 284 However, this would require follow-up of a large cohort (as incident TB is an uncommon event) 285 and could only be performed in groups who are not eligible for chemoprophylaxis.

To our knowledge, our study is the first evaluation of QFT-Plus assay among recent contacts of TB cases. Although limited by the small sample size, our data show that QFT-Plus in contact screening has an improved performance compared to QFT-GIT and suggests a role for the differential value between the two tubes as a proxy for recent infection. Larger prospective studies are needed to assess the positive predictive value of the test and the possible role of the differential value between the two antigens tube as marker for recent infection.

In conclusion, the difference between the two antigen tubes, used as an indirect estimate of specific CD8⁺ activation, is associated with factors indicating increased Mtb exposure, suggesting that this might identify individuals at greater risk of progression to active TB.

295 QFT-plus shows stronger association with surrogate measures of exposure compared to QFT-

GIT and therefore seems at least as accurate as QFT-GIT in the setting of contact screening.

297

298 ACKNOWLEDGMENTS

- 299
- 300 The authors thank the study subjects for their generous participation; the health care assistants
- and all the staff of the Region TB reference Centre Villa Marelli Niguarda Hospital for their
- 302 diligent work and valuable support. The authors thank QIAGEN (Hilden, Germany) for
- 303 providing QFT-Plus Kits free of charge.

REFERENCES

- 305
- H. Getahun, A. Matteelli, R. E. Chaisson, and M. Raviglione, "Latent Mycobacterium 306 [1] 307 tuberculosis Infection," N. Engl. J. Med., vol. 372, no. 22, pp. 2127–2135, May 2015. 308 [2] K. Lönnroth, G. B. Migliori, I. Abubakar, L. D'Ambrosio, G. de Vries, R. Diel, P. 309 Douglas, D. Falzon, M.-A. Gaudreau, D. Goletti, E. R. González Ochoa, P. LoBue, A. 310 Matteelli, H. Njoo, I. Solovic, A. Story, T. Tayeb, M. J. van der Werf, D. Weil, J.-P. 311 Zellweger, M. Abdel Aziz, M. R. M. Al Lawati, S. Aliberti, W. Arrazola de Oñate, D. 312 Barreira, V. Bhatia, F. Blasi, A. Bloom, J. Bruchfeld, F. Castelli, R. Centis, D. 313 Chemtob, D. M. Cirillo, A. Colorado, A. Dadu, U. R. Dahle, L. De Paoli, H. M. Dias, 314 R. Duarte, L. Fattorini, M. Gaga, H. Getahun, P. Glaziou, L. Goguadze, M. del 315 Granado, W. Haas, A. Järvinen, G.-Y. Kwon, D. Mosca, P. Nahid, N. Nishikiori, I. 316 Noguer, J. O'Donnell, A. Pace-Asciak, M. G. Pompa, G. G. Popescu, C. Robalo 317 Cordeiro, K. Rønning, M. Ruhwald, J.-P. Sculier, A. Simunović, A. Smith-Palmer, G. 318 Sotgiu, G. Sulis, C. A. Torres-Duque, K. Umeki, M. Uplekar, C. van Weezenbeek, T. 319 Vasankari, R. J. Vitillo, C. Voniatis, M. Wanlin, and M. C. Raviglione, "Towards 320 tuberculosis elimination: an action framework for low-incidence countries," Eur. 321 *Respir. J.*, Mar. 2015. 322 [3] R. Diel, R. Loddenkemper, J.-P. Zellweger, G. Sotgiu, L. D'Ambrosio, R. Centis, M. J. 323 van der Werf, M. Dara, A. Detjen, P. Gondrie, L. Reichman, F. Blasi, and G. B. 324 Migliori, "Old ideas to innovate TB control: preventive treatment to achieve 325 elimination," Eur. Respir. J., Feb. 2013. 326 C. Dye, S. Scheele, P. Dolin, V. Pathania, R. MC, and for the W. H. O. G. S. and M. [4] 327 Project, "Global burden of tuberculosis: Estimated incidence, prevalence, and mortality 328 by country," JAMA, vol. 282, no. 7, pp. 677–686, Aug. 1999. E. Vynnycky and P. E. M. Fine, "The natural history of tuberculosis: the implications 329 [5] 330 of age-dependent risks of disease and the role of reinfection," Epidemiol. Infect., vol. 331 119, no. 02, pp. 183–201, 1997. 332 [6] J. B. Bass, L. S. Farer, P. C. Hopewell, R. O'Brien, R. F. Jacobs, F. Ruben, D. E. 333 Snider, and G. Thornton, "Treatment of tuberculosis and tuberculosis infection in 334 adults and children. American Thoracic Society and The Centers for Disease Control 335 and Prevention.," Am. J. Respir. Crit. Care Med., vol. 149, no. 5, pp. 1359–1374, May 336 1994. 337 M. Farhat, C. Greenaway, M. Pai, And D. Menzies, "False-positive tuberculin skin [7]

- tests : what is the absolute effect of BCG and non-tuberculous mycobacteria?," *Int. J. Tuberc. lung Dis.*, vol. 10, no. 11, pp. 1192–1204.
- P. Andersen, M. E. Munk, J. M. Pollock, and T. M. Doherty, "Specific immune-based
 diagnosis of tuberculosis," *Lancet*, vol. 356, no. 9235, pp. 1099–1104, Jan. 2016.
- 342 [9] M. Sester, G. Sotgiu, C. Lange, C. Giehl, E. Girardi, G. B. Migliori, A. Bossink, K.
- 343 Dheda, R. Diel, J. Dominguez, M. Lipman, J. Nemeth, P. Ravn, S. Winkler, E. Huitric,
- A. Sandgren, and D. Manissero, "Interferon-γ release assays for the diagnosis of active
 tuberculosis: a systematic review and meta-analysis," *Eur. Respir. J.*, vol. 37, no. 1,
- 346 pp. 100–111, Jan. 2011.
- M. Santin, L. Muñoz, and D. Rigau, "Interferon-γ Release Assays for the Diagnosis of
 Tuberculosis and Tuberculosis Infection in HIV-Infected Adults: A Systematic Review
 and Meta-Analysis," *PLoS One*, vol. 7, no. 3, p. e32482, Mar. 2012.
- A. M. Mandalakas, A. K. Detjen, A. C. Hesseling, A. Benedetti, and D. Menzies,
 "Interferon-gamma release assays and childhood tuberculosis: systematic review and
 meta-analysis.," *Int. J. Tuberc. Lung Dis.*, vol. 15, no. 8, pp. 1018–32, Aug. 2011.
- M. X. Rangaka, K. A. Wilkinson, J. R. Glynn, D. Ling, D. Menzies, J. MwansaKambafwile, K. Fielding, R. J. Wilkinson, and M. Pai, "Predictive value of interferon-γ
 release assays for incident active tuberculosis: a systematic review and meta-analysis," *Lancet Infect. Dis.*, vol. 12, no. 1, pp. 45–55, Jun. 2015.
- V. Corbière, G. Pottier, F. Bonkain, K. Schepers, V. Verscheure, S. Lecher, T. M.
 Doherty, C. Locht, and F. Mascart, "Risk Stratification of Latent Tuberculosis Defined
 by Combined Interferon Gamma Release Assays," *PLoS One*, vol. 7, no. 8, p. e43285,
 Aug. 2012.
- 361 [14] G. Delogu, T. Chiacchio, V. Vanini, O. Butera, G. Cuzzi, A. Bua, P. Molicotti, S.
- Zanetti, F. N. Lauria, S. Grisetti, N. Magnavita, G. Fadda, E. Girardi, and D. Goletti,
 "Methylated HBHA produced in M. smegmatis discriminates between active and nonactive tuberculosis disease among RD1-responders.," *PLoS One*, vol. 6, no. 3, p.
 e18315, Jan. 2011.
- 366 [15] P. Escalante, T. Peikert, V. P. Van Keulen, C. L. Erskine, C. L. Bornhorst, B. R.
- 367 Andrist, K. McCoy, L. R. Pease, R. S. Abraham, K. L. Knutson, H. Kita, A. G.
- 368 Schrum, and A. H. Limper, "Combinatorial Immunoprofiling in Latent Tuberculosis
- Infection. Toward Better Risk Stratification," *Am. J. Respir. Crit. Care Med.*, vol. 192,
 no. 5, pp. 605–617, Jun. 2015.
- 371 [16] E. Petruccioli, L. Petrone, V. Vanini, G. Cuzzi, A. Navarra, G. Gualano, F. Palmieri, E.

372 Girardi, and D. Goletti, "Assessment of CD27 expression as a tool for active and latent 373 tuberculosis diagnosis," J. Infect., vol. 71, no. 5, pp. 526–533, Feb. 2016. 374 A. Rachow, N. Heinrich, and C. Geldmacher, "Early Identification of Progressive TB [17] 375 Disease Using Host Biomarkers," EBioMedicine, vol. 2, no. 2, pp. 107–108, Feb. 2016. 376 D. E. Zak, A. Penn-Nicholson, T. J. Scriba, E. Thompson, S. Suliman, L. M. Amon, H. [18] 377 Mahomed, M. Erasmus, W. Whatney, G. D. Hussey, D. Abrahams, F. Kafaar, T. 378 Hawkridge, S. Verver, E. J. Hughes, M. Ota, J. Sutherland, R. Howe, H. M. Dockrell, 379 W. H. Boom, B. Thiel, T. H. M. Ottenhoff, H. Mayanja-Kizza, A. C. Crampin, K. 380 Downing, M. Hatherill, J. Valvo, S. Shankar, S. K. Parida, S. H. E. Kaufmann, G. 381 Walzl, A. Aderem, and W. A. Hanekom, "A blood RNA signature for tuberculosis 382 disease risk: a prospective cohort study," Lancet, Apr. 2016. 383 [19] N. Rakotosamimanana, V. Richard, V. Raharimanga, B. Gicquel, T. M. Doherty, A. 384 Zumla, and V. Rasolofo Razanamparany, "Biomarkers for risk of developing active 385 tuberculosis in contacts of TB patients: a prospective cohort study," Eur. Respir. J., 386 Aug. 2015. 387 QUIAGEN, "QuantiFERON-TB-Gold-Plus ELISA Package Insert," 2014. [Online]. [20] 388 Available: www.QuantiFERON.com. 389 [21] D. A. Lewinsohn, A. S. Heinzel, J. M. Gardner, L. Zhu, M. R. Alderson, and D. M. 390 Lewinsohn, "Mycobacterium tuberculosis-specific CD8+ T Cells Preferentially Recognize Heavily Infected Cells," Am. J. Respir. Crit. Care Med., vol. 168, no. 11, 391 392 pp. 1346–1352, Dec. 2003. 393 C. L. Day, D. A. Abrahams, L. Lerumo, E. Janse van Rensburg, L. Stone, T. O'rie, B. [22] 394 Pienaar, M. de Kock, G. Kaplan, H. Mahomed, K. Dheda, and W. A. Hanekom, 395 "Functional Capacity of Mycobacterium tuberculosis-Specific T Cell Responses in 396 Humans Is Associated with Mycobacterial Load," J. Immunol., vol. 187, no. 5, pp. 397 2222-2232, Sep. 2011. 398 V. Rozot, S. Vigano, J. Mazza-Stalder, E. Idrizi, C. L. Day, M. Perreau, C. Lazor-[23] 399 Blanchet, E. Petruccioli, W. Hanekom, D. Goletti, P. A. Bart, L. Nicod, G. Pantaleo, 400 and A. Harari, "Mycobacterium tuberculosis-specific CD8+ T cells are functionally 401 and phenotypically different between latent infection and active disease," Eur. J. 402 Immunol., vol. 43, no. 6, pp. 1568–1577, 2013. 403 M. Nikolova, R. Markova, R. Drenska, M. Muhtarova, Y. Todorova, V. Dimitrov, H. [24] 404 Taskov, C. Saltini, and M. Amicosante, "Antigen-specific CD4- and CD8-positive 405 signatures in different phases of Mycobacterium tuberculosis infection," Diagn.

406		Microbiol. Infect. Dis., vol. 75, no. 3, pp. 277-281, 2013.
407	[25]	A. A. Ryan, J. K. Nambiar, T. M. Wozniak, B. Roediger, E. Shklovskaya, W. J.
408		Britton, B. Fazekas de St. Groth, and J. A. Triccas, "Antigen Load Governs the
409		Differential Priming of CD8 T Cells in Response to the Bacille Calmette Guérin
410		Vaccine or Mycobacterium tuberculosis Infection," J. Immunol., vol. 182, no. 11, pp.
411		7172–7177, Jun. 2009.
412	[26]	M. Faccini, S. Cantoni, G. Ciconali, M. T. Filipponi, G. Mainardi, A. F. Marino, S.
413		Senatore, L. R. Codecasa, M. Ferrarese, G. Gesu, E. Mazzola, And A. Filia,
414		"Tuberculosis-related stigma leading to an incomplete contact investigation in a low-
415		incidence country," Epidemiol. Infect., vol. 143, no. 13, pp. 2841-2848, 2015.
416	[27]	Tuberculosis. Clinical diagnosis and menegment of tuberculosis, and measures for its
417		prevention and control, no. March. 2011.
418	[28]	P. C. Hill, R. H. Brookes, A. Fox, K. Fielding, D. J. Jeffries, D. Jackson-Sillah, M. D.
419		Lugos, P. K. Owiafe, S. A. Donkor, A. S. Hammond, J. K. Otu, T. Corrah, R. A.
420		Adegbola, and K. P. W. J. McAdam, "Large-scale evaluation of enzyme-linked
421		immunospot assay and skin test for diagnosis of Mycobacterium tuberculosis infection
422		against a gradient of exposure in The Gambia.," Clin. Infect. Dis., vol. 38, no. 7, pp.
423		966–73, Apr. 2004.
424	[29]	J. Z. Metcalfe, A. Cattamanchi, C. E. McCulloch, J. D. Lew, N. P. Ha, and E. A.
425		Graviss, "Test Variability of the QuantiFERON-TB Gold In-Tube Assay in Clinical
426		Practice," Am. J. Respir. Crit. Care Med., vol. 187, no. 2, pp. 206-211, Jan. 2013.
427	[30]	R. Ribeiro-Rodrigues, S. Kim, F. D. Coelho da Silva, A. Uzelac, L. Collins, M. Palaci,
428		D. Alland, R. Dietze, J. J. Ellner, E. Jones-López, and P. Salgame, "Discordance of
429		Tuberculin Skin Test and Interferon Gamma Release Assay in Recently Exposed
430		Household Contacts of Pulmonary TB Cases in Brazil," PLoS One, vol. 9, no. 5, p.
431		e96564, May 2014.
432	[31]	S. W. Lee, D. K. Oh, S. H. Lee, H. Y. Kang, CT. Lee, and JJ. Yim, "Time interval
433		to conversion of interferon- γ release assay after exposure to tuberculosis," <i>Eur. Respir.</i>
434		J., vol. 37, no. 6, pp. 1447–1452, Jun. 2011.
435	[32]	M. Pai, K. Dheda, J. Cunningham, F. Scano, and R. O'Brien, "T-cell assays for the
436		diagnosis of latent tuberculosis infection: moving the research agenda forward," Lancet
437		Infect. Dis., vol. 7, no. 6, pp. 428-438, 2007.
438	[33]	T. Chiacchio, E. Petruccioli, V. Vanini, G. Cuzzi, C. Pinnetti, A. Sampaolesi, A.
439		Antinori, E. Girardi, and D. Goletti, "Polyfunctional T-cells and effector memory

- phenotype are associated with active TB in HIV-infected patients," *J. Infect.*, vol. 69,
 no. 6, pp. 533–545, 2014.
- 442 [34] C. Lancioni, M. Nyendak, S. Kiguli, S. Zalwango, T. Mori, H. Mayanja-Kizza, S.
- 443 Balyejusa, M. Null, J. Baseke, D. Mulindwa, L. Byrd, G. Swarbrick, C. Scott, D. F.
- Johnson, L. Malone, P. Mudido-Musoke, W. H. Boom, D. M. Lewinsohn, and D. a.
- 445 Lewinsohn, "CD8 + T cells provide an immunologic signature of tuberculosis in young
- 446 children," Am. J. Respir. Crit. Care Med., vol. 185, no. 2, pp. 206–212, 2012.
- 447 [35] L. Barcellini, E. Borroni, J. Brown, E. Brunetti, L. Codecasa, F. Cugnata, P. Dal
- 448 Monte, C. Di Serio, D. Goletti, G. Lombardi, M. Lipman, P. M. V Rancoita, M.
- Tadolini, and D. M. Cirillo, "First independent evaluation of QuantiFERON-TB Plus
- 450 performance," *Eur. Respir. J.*, Feb. 2016.
- 451
- 452

Table1: Demographic characteristics

	Number	%					
Sex (n=119)							
Male	63	52.9%					
Female	56	47.1%					
Estimated incidence of TB in country	y of birth $^{\circ}$ (n=	:119)					
0-50 per 100000 person-year	48	40.4%					
>50 per 100000 person-year	71	59.7%					
Country of birth (n=119)							
European	58	48.7%					
NON European	61	51.3					
BCG* vaccination (n=104)							
No	22	21.1%					
Yes	82	78.8%					
Smear status of index case (n=108)							
Negative	39	36.1%					
Positive	69	63.9%					
Time spent with the index case (hour	rs per day) (n=	=108)					
1-4	27	25%					
5-8	25	23.1%					
9-12	9	8.3%					
>12	47	43.5%					
Sleeping proximity to the index case	(n=108)						
Different house	61	56.5%					
Different rooms	19	17.6%					
Same room	28	25.9%					
Immunocompromised [§] (n=119)							
No	108	90.8%					
Yes	11	9.2%					

° As per WHO Report 2014 *BCG bacilli Clamette-Guérin

[§] Causes of immunosuppression: diabetes mellitus (6), chronic kidney disease (0), HIV (2), malignancy (2), immunosuppressive medications (1)

Table	2:	Test	results

QFT-GIT results	QFT Plu	Positive results per tube		QTF Plus IFN- γ concentrations (IU/ml)*			
results	Negative	Positive	TB1	TB2		TB1-Nil	TB2-Nil
Negative (n=63)	51 (80.95%)	12 (19.05%)	10°	10 [§]		0.01 (-0.01;0.17)	0.04 (0;0.23)
Positive (n=56)	0	56 (100%)	56	56		10.60 (2.94;16.57)	11.00 (3.32;17.75)
Total (n=119)	51 (42.86%)	68 (57.14%)	66	66		0.74 (0.01;9.65)	0.67 (0.04;8.94)

*median (25-75 percentile) ° 2 were positive to TB1 only § 2 were positive to TB2 only

Sample no	BCG scar	TST**	QFT- GIT	QFT- Plus TB1°	QFT- Plus TB2 [§]	Index case smear status	Relation to Index case	Immunosuppression
C1	Yes	20	Neg	1.83	0.51	Pos	Household, primary caregiver	Prednisone treatment
C11	Yes	7	Neg*	0.49	0.83	Pos	Boyfriend	No
C15	No	21	Neg*	0.11	0.48	Pos	Employer (index case: house-made)	No
C17	Yes	10	Neg	0.38	0.41	Neg	Household, sister	No
C39	Yes	20	Neg	0.83	0.88	Pos	Hospital close contact (sharing the same room)	Cancer
C53	Yes	20	Neg	0.3	0.58	Pos	Colleague, every day ride at work	No
C63	Yes	16	Neg	0.74	0.67	Neg	Household	No
C69	Yes	14	Neg	0.52	0.29	Pos	Household	No
C75	Yes	11	Neg	0.81	0.9	Pos	Household	No
C78	No	11	Neg	1.88	1.93	Pos	Colleague (sharing the same room)	No
C91	Yes	14	Neg	0.36	0.1	Neg	Household	No
C98	Yes	20	Neg	1.65	1.14	Pos	Household	Pregnant

Table 3: QFT-Plus and QFT-GIT discordant results

°TB1-Nil

⁸ TB2-Nil
* Repeated test by QFT-IT at follow up converted to positive
** Diameter of induration in mm

				•,•		0.6
	QFT Positive		QFT Plus Pos		TB2-TB1>0.6	
	OR (95% CI)	p-value	OR (95% CI)	p-value	OR (95% CI)	p-value
Age	1.02 (1.00;1.05)	0.09	1.02 (1.00;1.05)	0.101	1.03 (1.00;1.07)	0.053
Sex						
Male	1		1		1	
Female	0.73 (0.35;1.5)	0.387	0.76 (0.37;1.57)	0.458	1.49 (0.55;4.10)	0.435
Estimated incidence	of TB per 100000 p	erson-year	in country of birth°			
0-50	1		1		1	
>50	1.44 (0.69;3.01)	0.333	1.63 (0.78;3.42)	0.197	0.82 (0.30;2.25)	0.7
Country of birth						
NON European	1		1		1	
European	0.84(0.41;1.73)	0.635	0.74 (0.36;1.54)	0.428	3.24 (1.07;9.75)	0.037
BCG* vaccination						
No	1		1		1	
Yes	2.04 (0.75;5.53)	0.161	2.26 (0.87;5.89)	0.096	0.98 (0.25;3.87)	0.978
Smear status of index	x case					
Negative	1		1		1	
Positive	1.12 (0.51;2.47)	0.780	1.39 (0.63;3.07)	0.413	1.29 (0.41;4.03)	0.662
Time spent with the	index case (hours pe	er day)				
1-4	1		1		1	
5-8	1.65 (0.48;5.67)	0429	3.23 (0.97;10.72)	0.055	3.55 (0.34;36.53)	0.288
9-12	2.8 (0.57;13.83)	0.206	4.37 (0.89;21.61)	0.070	7.43 (0.59;94.26)	0.122
>12	6.78 (2.28;20.16)	0.0006	14.78(4.62;47.25)	5.6e-06	7.03 (0.85;58.2)	0.071
Sleeping proximity to	o the index case					
Different house	1		1		1	
Different rooms	3.79 (1.29;11.14)	0.015	5.78 (1.71;19.52)	0.005	0.51 (0.06;4.52)	0.545
Same room	3.98 (1.55;10.23)	0.004	5.65 (2.00;15.97)	0.001	4.34 (1.37;13.81)	0.013
Immunocompromised						
No	1		1		1	
Yes	0.62 (0.17;2.22)	0.459	1.35 (0.37;4.88)	0.649	0.54 (0.06;4.46)	0.563
° As per WHO Repor			,		,	

*BCG bacilli Clamette-Guérin

Table 5. Backward stepwise multivariate logistic regressions for predicting QFT-GIT or **QFT Plus Positivity**

	QFT-GIT Pos	QFT-GIT Positive OR (95% CI) p-value		itive			
	OR (95% CI)			p-value			
Time spent with the index case (hours per day)							
1-12	1		1				
>12	4.63 (2.05; 10.47)	0.0002	6.98 (2.86; 17.02)	1.98e-05			
0.4 0.4 0.14							

° As per WHO Report 2014

	OR (95% CI)	p-value
Country of birth		
NON European	1	
European	3.46 (1.03;11.69)	0.0453
Sleeping proximity to the	index case	
No same room	1	
Same room	5.90 (1.83;18.97)	0.0029

Table 6. Backward stepwise multivariate logistic regression for predicting TB2-TB1>0.6