



Biomarkers to identify *Mycobacterium tuberculosis* infection among borderline QuantiFERON results

Jonathan W. Uzorka¹, Jaap A. Bakker ², Krista E. van Meijgaarden¹, Eliane M.S. Leyten³, Nathalie M. Delfos⁴, David J. Hetem⁵, Jos Kerremans⁶, Mieke Zwarts⁷, Sandra Cozijn⁸, Tom H.M. Ottenhoff¹, Simone A. Joosten¹ and Sandra M. Arend¹

¹Dept of Infectious Diseases, Leiden University Medical Centre, Leiden, The Netherlands. ²Dept of Clinical Chemistry and Laboratory Medicine, Leiden University Medical Centre, Leiden, The Netherlands. ³Dept of Internal Medicine, Haaglanden Medical Centre, Den Haag, The Netherlands. ⁴Dept of Internal Medicine, Alrijne Hospital, Leiderdorp, The Netherlands. ⁵Dept of Medical Microbiology, Haaglanden Medical Centre, Den Haag, The Netherlands. ⁶Dept of Medical Microbiology, Alrijne Hospital, Leiderdorp, The Netherlands. ⁷Dept of Clinical Chemistry and Laboratory Medicine, Haaglanden Medical Centre, Den Haag, The Netherlands. ⁸Dept of Medical Microbiology, Alrijne Hospital, Leiden, The Netherlands.

Corresponding author: Jonathan W. Uzorka (j.w.uzorka@lumc.nl)



Shareable abstract (@ERSpublications) Additional laboratory testing of IP-10/CXCL10 and MIG/CXCL9 in individuals at increased risk of reactivation TB with a borderline QuantiFERON-TB Gold Plus result just below the formal cut-off helps to identify those with *M. tuberculosis* infection https://bit.ly/3t8zR7r

Cite this article as: Uzorka JW, Bakker JA, van Meijgaarden KE, *et al.* Biomarkers to identify *Mycobacterium tuberculosis* infection among borderline QuantiFERON results. *Eur Respir J* 2022; 60: 2102665 [DOI: 10.1183/13993003.02665-2021].

This single-page version can be shared freely online.

Copyright ©The authors 2022.

Abstract

This version is distributed under the terms of the Creative Commons Attribution Non-Commercial Licence 4.0. For commercial reproduction rights and permissions contact permissions@ersnet.org

This article has an editorial commentary: https://doi.org/10.1183/ 13993003.00697-2022

Received: 8 Oct 2021 Accepted: 18 Dec 2021 **Background** Screening for tuberculosis (TB) infection often includes QuantiFERON-TB Gold Plus (QFT) testing. Previous studies showed that two-thirds of patients with negative QFT results just below the cutoff, so-called borderline test results, nevertheless had other evidence of TB infection. This study aimed to identify a biomarker profile by which borderline QFT results due to TB infection can be distinguished from random test variation.

Methods QFT supernatants of patients with a borderline (≥ 0.15 and $< 0.35 \text{ IU} \cdot \text{mL}^{-1}$), low-negative ($< 0.15 \text{ IU} \cdot \text{mL}^{-1}$) or positive ($\ge 0.35 \text{ IU} \cdot \text{mL}^{-1}$) QFT result were collected in three hospitals. Bead-based multiplex assays were used to analyse 48 different cytokines, chemokines and growth factors. A prediction model was derived using LASSO regression and applied further to discriminate QFT-positive *Mycobacterium tuberculosis*-infected patients from borderline QFT patients and QFT-negative patients

Results QFT samples of 195 patients were collected and analysed. Global testing revealed that the levels of 10 kDa interferon (IFN)- γ -induced protein (IP-10/CXCL10), monokine induced by IFN- γ (MIG/CXCL9) and interleukin-1 receptor antagonist in the antigen-stimulated tubes were each significantly higher in patients with a positive QFT result compared with low-negative QFT individuals (p<0.001). A prediction model based on IP-10 and MIG proved highly accurate in discriminating patients with a positive QFT (TB infection) from uninfected individuals with a low-negative QFT (sensitivity 1.00 (95% CI 0.79–1.00) and specificity 0.95 (95% CI 0.74–1.00)). This same model predicted TB infection in 68% of 87 patients with a borderline QFT result.

Conclusions This study was able to classify borderline QFT results as likely infection-related or random. These findings support additional laboratory testing for either IP-10 or MIG following a borderline QFT result for individuals at increased risk of reactivation TB.

