

FUNCTIONAL, ANTIGEN-SPECIFIC STEM CELL-LIKE MEMORY (T_{SCM}) CD4⁺ T CELLS ARE INDUCED BY HUMAN *M. TUBERCULOSIS* INFECTION

Cheleka Mpande, One B. Dintwe, Munya Musvosvi, Mark Hatherill, Elisa Nemes and Thomas J. Scriba

South African Tuberculosis Vaccine Initiative, University of Cape Town, South Africa.

Background

Induction of anti-mycobacterial T cell responses by vaccination or natural infection is not sufficient to consistently confer protection against TB disease. A better understanding of the *Mycobacterium tuberculosis* (M.tb)-specific memory T cell repertoire is needed. Compared to central (T_{CM}) and effector (T_{EFF}) memory T cells, stem cell-like memory T cells (T_{SCM}) have superior self-renewing capacity, longevity and proliferative potential, functions associated with maintenance of long lasting immunity. Our knowledge of T_{SCM} function derives primarily from studies of virus-specific CD8⁺ T_{SCM}. Little is known about CD4⁺ T_{SCM} function in response to bacterial infections. We aimed to determine if M.tb infection generates antigen-specific CD4⁺ T_{SCM} and if so, characterise their functional ontology.

Methods

We studied immune responses to natural M.tb infection in PBMCs from three cross-sectional QuantiFERON (QFT)+ adult cohorts and a longitudinal adolescent cohort of recent QFT converters. M.tb-specific CD4 T cells were detected by flow cytometry using MHC-class II tetramers bearing Ag85, CFP-10 or ESAT-6 peptides, or by intracellular cytokine staining after whole blood stimulation with Ag85B, CFP-10, ESAT-6 or BCG. Transcriptional analyses of M.tb-specific tetramer+ CD4⁺ (Tet⁺) T_{SCM} (CD45RA+CCR7+CD27+), sorted by FACS, were performed by microfluidic quantitative real-time PCR. The transcriptomic profile of Tet⁺ T_{SCM} was confirmed on independent cohorts by measuring expression of chemokine receptors, cytotoxic molecules and cytokines using flow cytometry.

Results

M.tb-specific Tet⁺ T_{SCM} were induced by primary M.tb-infection and maintained thereafter. CD95⁻ Tet⁺ T_{SCM} displayed a very distinct transcriptomic profile from bulk CD4⁺ naïve T cells (T_N) and more similar to bulk T_{SCM}, M.tb-specific T_{SCM} and T_{EFF}. CD95 expression, a typical marker of CD8⁺ T_{SCM}, by Tet⁺ T_{SCM} was variable. By contrast, Tet⁺ T_{SCM} were predominantly CXCR3⁺, an alternative T_{SCM} marker. Tet⁺ T_{SCM} expressed significantly higher protein levels of CCR5, CCR6, CXCR3, Granzyme A, Granzyme K and Granulysin than bulk T_N and CD95⁺ T_{SCM}. These M.tb-specific T_{SCM} were also functional, producing IL-2, IFN-γ and TNF-α upon antigen stimulation.

Conclusions

Our comprehensive analyses show that M.tb CD4⁺ T_{SCM} are induced during primary M.tb-infection and are distinct from bulk T_N and T_{SCM}. CXCR3 was a more robust marker of M.tb-specific CD4⁺ T_{SCM} than CD95. M.tb-specific CD4⁺ T_{SCM} displayed chemokine receptor profiles consistent with memory Th1/17 cells and, surprisingly, expressed effector functions including cytotoxic markers and Th1 cytokines.

Funding: South African Medical Research Council