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# Altered monocyte phenotypes but not impaired peripheral T cell immunity may explain susceptibility of the elderly to develop tuberculosis

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# Abstract

Tuberculosis (TB) is the leading killer due to a single infectious disease worldwide. With the aging of the global population, the case rate and deaths due to TB are highest in the elderly population. While general immunosenescence associated with old age is thought to contribute to the susceptibility of the elderly to develop active TB disease, very few studies of immune function in elderly individuals with Mycobacterium tuberculosis (M.tb) infection or disease have been performed. In particular, impaired adaptive T cell immunity to *M.tb* is one proposed mechanism for the elderly's increased susceptibility primarily on the basis of the decreased delayed type hypersensitivity response to tuberculin-purified protein derivative in the skin of elderly individuals. To investigate immunological reasons why the elderly are susceptible to develop active TB disease, we performed a cross-sectional observational study over a five year period (2012–2016) enrolling participants from 2 age groups (adults: 25–44 years; elderly: 65 and older) and 3 *M.tb* infection statuses (active TB, latent TB infection, and healthy controls without history of *M.tb* infection). We hypothesized that impaired peripheral T cell immunity plays a role in the biological susceptibility of the elderly to TB. Contrary to our hypothesis, we observed no evidence of impaired *M.tb* specific T cell frequency or altered production of cytokines implicated in *M.tb* control (IFN- $\gamma$ , IL-10) in peripheral blood in the elderly. Instead, we observed alterations in monocyte proportion and phenotype with age and *M.tb* infection that suggest their potential role in

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**Declarations of interest** 

None.

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the susceptibility of the elderly to develop active TB. Our results suggest a potential link between the known widespread low-grade systemic inflammation of old age, termed "inflammaging," with the elderly's specific susceptibility to developing active TB. Moreover, our results highlight the need for further research into the biological reasons why the elderly are more susceptible to disease and death from TB, so that public health systems can be better equipped to face the present and future problem of TB in an aging global population.

## **Keywords**

Tuberculosis; Geriatrics; T cell; Monocyte; Inflammaging; Immunosenescence

# 1. Introduction

The global population is aging. It is estimated that by the year 2050, 17% of the global population will be 65 years of age or older, compared to 8.5% in 2015 (US Census Bureau, 2016). Increasing age is the major risk factor for a host of diseases including diabetes, cardiovascular disease, cancer and neurodegenerative disease (López-Otín et al., 2013; Niccoli and Partridge, 2012). Increasing age is also a major risk factor for infectious diseases (Gavazzi and Krause, 2002; Yoshikawa, 1997). Tuberculosis (TB), caused by the airborne pathogen *Mycobacterium tuberculosis* (*M.tb*), is currently the leading killer due to an infectious disease, and increasing age is a significant risk factor for disease and death due to TB. For example, in the United States, the case rate of symptomatic, active TB is highest in those 65 years of age, and this age group is the most likely to die from TB (Centers for Disease Control and Prevention [CDC], 2017). Worldwide, the case rate is similarly highest among those 55 years of age (World Health Organization [WHO], 2017). Moreover, according to the 2016 Global Burden of Disease estimate, 62% of deaths due to TB globally occurred among people older than 50, with more than half of these deaths occurring in those

65 years of age (Institute for Health Metrics and Evaluation, 2016). We can expect that as the global population continues to age, the global burden of TB in the elderly will also increase (Negin et al., 2015).

Infection with *M.tb* is thought to confer a ~10% lifetime risk of developing symptomatic, transmissible active TB, with the remainder of individuals controlling the bacteria in a state known as latent TB infection (LTBI) (Vynnycky and Fine, 2000). While the prevalence of active TB has decreased in the last 50 years, the increased risk of the elderly to convert from LTBI to active TB poses an additional challenge to TB elimination efforts in the foreseeable future (Glaziou et al., 2015; Hochberg and Horsburgh, 2013; Horsburgh et al., 2010; Vynnycky and Fine, 2000). Research into the clinical and biological reasons for why the elderly are especially susceptible to develop active TB can help inform strategies to combat this present and future public health problem.

General immunosenescence associated with old age has been cited as one biological reason for the increased susceptibility of the elderly to develop active TB (Byng-Maddick and Noursadeghi, 2016; Guzzetta and Kirschner, 2013; Menon et al., 2016; Rajagopalan, 2016; Thrupp et al., 2004). However, very few studies of immune function in elderly individuals

with *M.tb* infection (LTBI or active TB) have been performed. A historical hypothesis in the field is that impaired adaptive T cell immunity to *M.tb* plays a role in this biological susceptibility. This hypothesis stems from observations that reactivity to tuberculin-purified protein derivative (PPD) in the tuberculin skin test (TST), a measurement of a delayed type hypersensitivity (DTH) response for LTBI diagnosis, is reduced with advancing age past 65 years despite epidemiological evidence of higher exposure earlier in life (Dorken et al., 1987; Van den Brande and Demedts, 1992). A few articles have shown that elderly patients with active TB had increased T-suppressor cell activity, or suppression of lymphocyte proliferation *in vitro* (Adambekov and Morozov, 1989; Harada, 1989). More recently, T cell responses to specific latency-associated antigens of *M.tb* have been investigated in the elderly, though without specific comparison to younger cohorts (Commandeur et al., 2011). These studies, taken together, do not provide definitive proof that age-related changes in T cell function contribute to risk of active TB in the elderly.

To provide additional investigation of immunological reasons why the elderly are susceptible to develop active TB, we performed a cross-sectional observational study over a five year period (2012–2016) enrolling participants from 2 age groups (adults: 25–44 years; elderly: 65 and older) and 3 *M.tb* infection statuses (active TB, LTBI, and healthy controls without history of *M.tb* infection). We hypothesized that impaired peripheral T cell immunity plays a role in the biological susceptibility of the elderly to TB. Contrary to our hypothesis, we observed no evidence of impaired *M.tb* specific T cell frequency or altered production of cytokines important for *M.tb* control in peripheral blood in the elderly. Instead, we observed alterations in monocyte proportion and phenotype with age and *M.tb* infection that suggest their potential role in the susceptibility of the elderly to develop active TB. Our results suggest a potential link between the known widespread low-grade systemic inflammation of old age, termed "inflammaging," with the elderly's specific susceptibility to developing active TB (Franceschi and Campisi, 2014; Piergallini and Turner, 2017).

# 2. Materials and Methods

# 2.1 Subject Recruitment

Study participants were from two age groups, 25–44 years and 65 years and older. Subjects were recruited from The Ohio State University Wexner Medical Center and Columbus Public Health. Eligible subjects were enrolled in the inpatient and outpatient setting and consented according to an IRB approved protocol. Subjects' medical histories were obtained by a questionnaire at study enrollment, and this information was verified and additional information obtained by review of their medical records. Clinical data collected for active TB/LTBI diagnosis included signs and symptoms of active TB, physical examination, QuantiFERON®TB-Gold Test (QFT) (Qiagen, USA), acid fast bacilli smears, *M.tb* PCR, cultures and imaging studies if available. Most subjects with active TB/LTBI were recruited at or before the day they started receiving treatment for active TB/LTBI. Upon study enrollment, peripheral blood was collected in lithium heparin coated tubes (Becton, Dickinson, NJ). Time from blood collection to peripheral blood mononuclear cell (PBMC) isolation was 4 hours.

# 2.2 PBMC Isolation

PBMCs were isolated by Ficoll-Paque (GE Healthcare Life Sciences) density centrifugation and washed twice before use in subsequent assays.

# 2.3 Measurement of ESAT-6 and CFP-10 specific IFN-γ+ T cells

The T-SPOT<sup>®</sup>.TB (Oxford Immunotec, UK) ELISPOT kit was used to measure the frequency of ESAT-6 and CFP-10 specific IFN- $\gamma$ + T cells in PBMCs, per the manufacturer's protocol. Spots were counted manually by two lab members independently. The results gave identical trends between counters (results of one count are shown). In accordance with the manufacturer's protocol, the data shown are from counts with the count of the negative control well subtracted.

# 2.4 PBMC Culture and Stimulation

 $2.5 \times 10^5$  PBMCs were cultured in triplicate in AIM-V media (with Albumax, 50 µg/mL Streptomycin, 10 µg/mL Gentamycin) (GIBCO) with 10 µg/mL H37Rv culture filtrate protein (CFP) (BEI Resources) or either media only or 10 µg/mL concanavalin A (Sigma-Aldrich) as negative/positive controls, respectively. Cell-free supernatants were removed after 48 hours culture at 37°C, 5% CO<sub>2</sub> and stored at -20 °C until further analysis.

# 2.5 Measurement of Cytokine Levels

Human IFN- $\gamma$ , IL-10 and IL-4 were measured in PBMC culture supernatants using ELISA kits (BD Biosciences), according to manufacturer's instructions. Samples outside the linear range were serially diluted and re-assayed. Some samples were thawed twice. IFN- $\gamma$  data from the first 72 recruited subjects were excluded from analysis due to the assay falling outside the linear range of the standards and samples being insufficient for further testing.

# 2.6 Flow Cytometry

 $5 \times 10^5$  PBMCs were washed in flow buffer (RPMI-1640, 10.4 mM HEPES, 0.1% sodium azide), stained for 20 minutes at 4°C with antibody and washed twice more. Antibodies used were: CD4-APC-Cy7 (Clone: RPA-T4), CD8-PE (Clone: HIT8a), CCR7-PerCP-Cy5.5 (Clone: 150503), PD-1-APC (Clone: MIH4), CD16-FITC (Clone: 3G8) and CD14-PerCP-Cy5.5 (Clone: MΦ9). All antibodies were purchased from BD Pharmingen. Data were acquired on a BD FACS Canto II with FACS Diva software (BD Biosciences). 50,000 events per sample were acquired. The data were analyzed using FlowJo software (Version 10.3.0). Lymphocytes were identified by FSC/SSC before gating on CD4 and CD8. The CD4/CD8 ratio was calculated from these gates. CCR7 and PD-1 positivity were determined using isotype controls. Monocytes were identified by gating on PBMCs using FSC/SSC before gating on CD14+ PBMCs. Monocyte subsets were subsequently identified by CD14 and CD16: classical (CD14++CD16-), intermediate (CD14++CD16+) and nonclassical (CD14dimCD16+).

## 2.7 Statistical Analysis

Data were analyzed using R v. 3.4.1 (R Foundation for Statistical Computing, Vienna, AUT). The chi-squared test or ANOVA were used to assess univariate differences in clinical

covariates between our main study groups, for categorical and continuous variables, respectively. The non-parametric Mann-Whitney U-test was used in all pair-wise comparisons for T cell function assay data. For our T cell and monocyte flow cytometry data, the non-parametric Kruskal-Wallis test was used, after which Dunn's *post hoc* test was used for pair-wise comparisons if the Kruskal-Wallis test gave p < 0.05. To assess the contribution of confounding clinical covariates to these comparisons, multivariate linear models were fit using the "Im" function in R and analysis of variance performed using the "ANOVA" function. IFN- $\gamma$  ELISA data were log-transformed before performing multivariate analysis.

# 3. Results

## 3.1 Cohort Description

We recruited a total of 205 subjects to our study, and a total of 169 subjects were included in the final analysis. Subjects excluded from analysis mostly included those whose samples were used for assay standardization or who had a previous history of treatment for active TB or LTBI (see Figure S1). 6 study groups were defined for our primary analysis, with 2 age groups (adults: 25–44 years; elderly: 65 and older) and 3 *M.tb* infection statuses (active TB, LTBI, healthy controls without history of *M.tb* infection). All but 3 subjects with active TB were culture confirmed to have *M.tb* complex, with the other three diagnoses being clinical TB uveitis disease which improved on TB treatment and two subjects with pleural TB diagnosed based on abnormal chest x-ray and chest-computed tomography (CT) and improvement following TB treatment. Subjects with LTBI were either clinically diagnosed by a positive QFT (n=39), or had a positive TST (n=1). All were deemed absent of active TB based on symptoms, chest x-ray, CT and/or negative sputum samples. All control subjects had a negative QFT result (n=18) or no known QFT result with no prior history of a positive TST (n=98). No control subjects had a clinical history of active TB. Moreover, most control subjects originated from a location of very low TB incidence.

Table 1 shows clinical covariates for our study subjects, and Table 2 shows culture confirmation and tissue site affected in our subjects with active TB. Due to our complex study design and available patient populations, not all covariates are balanced across all 6 study groups. For several of our immunological assays showing significant differences between age groups, we used multivariate analysis to assess the effects of our known clinical confounding variables.

# 3.2 T cell cytokine responses to *M.tb* antigens are not altered by age in those with LTBI or active TB

We first sought to determine whether among *M.tb* infected individuals, the elderly had decreased peripheral *M.tb* specific T cell numbers and function compared to adults. We measured the frequency of ESAT-6 and CFP-10 specific T cells using the T-SPOT.TB ELISPOT kit. Of note, peptide lengths (15-mer) were designed to contain MHC II epitopes (Li et al., 2012; Richeldi, 2006). We observed no significant differences in numbers of both ESAT-6 and CFP-10 specific IFN- $\gamma$ + T cells in the elderly relative to adults, in both latent and active TB groups (p>0.05, see Figure 1A–B). In fact, we observed a trend for increased

ESAT-6 and CFP-10 specific IFN- $\gamma$ + T cells in elderly subjects with LTBI compared to adults. It is possible that while the elderly have increased CFP-10 and ESAT-6 specific T cells in the blood, the amount of IFN- $\gamma$  that these and other cells produce is decreased. Indeed, the ability of T cells to produce IFN- $\gamma$  in response to the large collection of *M.tb* antigens contained in *M.tb* culture filtrate protein (CFP) is decreased in adult active TB cases with higher clinical disease severity (Jamil et al., 2007). To test whether *M.tb*-specific IFN- $\gamma$  production was reduced in the elderly, we stimulated PBMCs with *M.tb* CFP and measured supernatant IFN- $\gamma$  by ELISA. Contrary to our hypothesis, there was no significant difference in CFP-specific IFN- $\gamma$  production with age, either among subjects with LTBI or active TB (p>0.05, see Figure 1C). We also measured IL-10 and IL-4 production by CFPstimulated PBMCs. We observed a trend for an increase in IL-10 production by CFPstimulated PBMCs from subjects with active TB relative to those with latent TB, consistent with what is known in the TB literature (Beamer et al., 2008a; Bonecini-Almeida et al., 2004; Jamil et al., 2007) (Figure 1D). However, we saw no significant differences in IL-10 production between age groups among those with LTBI or active TB (p>0.05). Moreover, we observed no change in IL-4 production between age groups, neither basally nor with CFP-stimulation (data not shown). Taken together, our data on M.tb-specific T cell frequency and cytokine production provide no evidence of *M.tb*-specific T cell dysfunction in the periphery in elderly people 65 years of age.

Interestingly, we did observe a statistically significant decrease in IFN- $\gamma$  production in response to CFP in the elderly who had no history of *M.tb* infection, relative to adults (Figure 1C). This suggests potential hypo-responsiveness of the innate immune system to mycobacterial antigens in the elderly. This observation was not confounded by diabetes, cancer, immunosuppressive treatment or the presence of another comorbidity, using graphical analysis (data not shown). Accordingly, the effect of age alone remained significant using a multivariate linear model including these variables (p=0.007) or all categorical clinical covariates from Table 1 (p=0.005). In addition, BCG vaccination status had no effect (p>0.05).

# 3.3 Global T cell phenotypes are altered by *M.tb* infection and disease independent of age

We also studied global T cell phenotypes by flow cytometry in our cohort. Consistent with what is known in the clinical literature, the CD4/CD8 ratio was decreased in subjects with LTBI and was further decreased in subjects with active TB relative to controls (data not shown). There was no significant difference in the CD4/CD8 ratio with age in any group (data not shown). We further sought to determine whether global T cell memory phenotype or exhaustion status was altered by *M.tb* infection and age. We observed a trend for a decrease in both CD4+CCR7+ and CD8+CCR7+ lymphocytes in adults with LTBI and a trend for a decrease in adults with active TB, with significance reached for CD8+CCR7+ lymphocytes (p<0.05), relative to adult controls (Figure 2A–B). However, we observed no significant difference in these parameters between the elderly and adults among those with LTBI or active TB (Figure 2A–B). Among our controls without history of *M.tb* infection, we did observe a significant decrease in CD8+CCR7+ lymphocytes and a trend for decreased CD4+CCR7+ lymphocytes with age, consistent with the aging literature (Koch et al., 2008; Yan et al., 2010) (Figures 2A–B). As for markers of T cell exhaustion, in a majority of

subjects we measured no PD-1+ CD4+ or CD8+ lymphocytes (Figures 2C–D). There was a trend for increased PD1+ cells in both CD4+ and CD8+ lymphocytes in elderly with active TB relative to all other groups, though our low sample size in this group makes this conclusion tentative. Taken together with our cytokine production data, our data do not suggest a clear mechanism for impaired peripheral T cell immunity as a dominant cause of potential biological susceptibility of the elderly to active TB.

# 3.4 Old age and *M.tb* infection alter the monocyte/lymphocyte ratio and skew monocytes towards a nonclassical phenotype

It is increasingly appreciated that monocyte phenotypes correlate with susceptibility to a variety of diseases (Berry et al., 2010; Puchta et al., 2016; Rogacev et al., 2012; Shaw et al., 2010; Wong et al., 2012). In addition to our examination of T cell phenotype and function, we sought to determine whether monocyte proportions and phenotypes were altered by old age and *M.tb* infection. In children and adults, the monocyte/lymphocyte ratio is associated with risk of active TB and disease severity, with a higher ratio being associated with highest risk and a very low ratio also associated with increased risk (Naranbhai et al., 2014a, 2014b). Interestingly, we observed an increased monocyte/lymphocyte ratio in the elderly with active TB, relative to all groups (p<0.05) (Figure 3A). We also observed a decreased monocyte/lymphocyte ratio in the elderly with LTBI, relative to both control groups and adults with LTBI (p<0.05 for controls and p=0.062 for adults with LTBI). In our examination of monocyte subset proportions defined by CD14 and CD16 expression, we observed a significant increase in the proportion of nonclassical (CD14dimCD16+) monocytes in elderly controls relative to adults, which has been observed by others (Hearps et al., 2012; Nyugen et al., 2010; Pence and Yarbro, 2018) (Figure 3B). This increase remained significant when taking into account diabetes, cancer, immunosuppressive treatment and the presence of another comorbidity (p<0.001) or all categorical clinical covariates from Table 1 (p<0.001) (see Table 3). Even more interesting, we observed a marginally significant increase in nonclassical monocytes in the elderly with LTBI, compared to elderly controls and adults with LTBI (p=0.106 and p<0.05, respectively). This suggests the hypothesis of an age-dependent increase in nonclassical monocytes in LTBI. Mirroring the results in nonclassical monocytes, we observed a trend for a decreased classical monocyte (CD14++CD16-) proportion in elderly controls relative to adults and a trend for a decreased proportion in elderly with LTBI relative to elderly controls and adults with LTBI (data not shown). We observed no significant differences in intermediate (CD14+ +CD16+) monocyte proportions with age or *M.tb* infection status (data not shown). Contrary to our results examining *M.tb*-specific T cell function in the elderly, our monocyte data suggest the hypothesis that changes in monocyte proportion and phenotype may contribute to susceptibility to develop active TB in the elderly.

# 4. Discussion

The global burden of TB in the elderly will increase as the global population continues to age (Negin et al., 2015). From available epidemiologic data (CDC, 2017; Negin et al., 2015; WHO, 2017), it is clear that changes incident to age contribute to the likelihood of the elderly to die from an episode of active TB. In areas of low TB incidence, the increased

likelihood of the elderly to develop active TB could be due to two effects. Epidemiologically, the elderly had higher exposure to TB in earlier years of life, when TB was more prevalent, compared to younger birth cohorts. Thus, a higher case rate in the elderly would simply reflect a larger reservoir of LTBI that could reactivate to active TB (Powell and Farer, 1980; Vynnycky and Fine, 2000; Wiker et al., 2010). Conversely, the increased case rate of TB in the elderly could be due to an age-related biological susceptibility to transition from a latent TB infection to active TB (Hochberg and Horsburgh, 2013; Horsburgh et al., 2010; Tocque et al., 1998). In South-East Asia, where new and relapse TB case notification rates are in general on par with case rates in Africa, those older than 55 years of age still have the highest incidence rate of TB, suggesting that the elderly are biologically susceptible to developing active TB (Tocque et al., 1998; WHO, 2017; Yew Wing W. et al., 2018). Our cross-sectional observational study assumes that the elderly are biologically more susceptible to developing active TB.

We hypothesized that impaired T cell immunity to *M.tb* plays a role in the biological susceptibility of the elderly to develop active TB. This has been a historical hypothesis in the field, stemming mainly from observations that reactivity to the PPD skin test, a measurement of a DTH response for LTBI diagnosis, is reduced with advancing age past 65 years despite epidemiological evidence of higher exposure earlier in life (Dorken et al., 1987; Van den Brande and Demedts, 1992). Contrary to our hypothesis, we observed no difference between adults and elderly people with LTBI or active TB when we measured the frequency of IFN- $\gamma$ producing T cells specific to ESAT-6 or CFP-10 or the amount of IFN- $\gamma$  or IL-10 produced in response to *M.tb* culture filtrate protein in PBMCs. Similarly, we saw no statistically significant effect of age on CCR7+ subsets of CD4 or CD8 T cells among those with LTBI or active TB. Therefore, among these measurements of *M.tb* specific and global T cell phenotype and function, we saw no evidence of impaired peripheral T cell immunity as a dominant cause of biological susceptibility of the elderly to develop active TB. The discordance of our functional T cell data with these historical PPD results in the elderly is consistent with the work of Mori and Kobashi which demonstrated that though PPD positive results are dramatically reduced in patients over the age of 80 who have culture-positive TB, their reactivity to a CFP-10 and ESAT-6 IFN- $\gamma$  release assay using peripheral blood is not diminished relative to younger adults (Kobashi et al., 2008; Mori et al., 2004). It is important to note that ESAT-6 and CFP-10 are immunodominant *M.tb* antigens and are not present in BCG or most nontuberculous mycobacteria (NTM), including *M. avium*. We chose *M.tb* culture filtrate protein, which does contain many epitopes in NTM and BCG, to measure total cytokine production in subject's PBMCs to assess the wider antigenic response to *M.tb.* Importantly, the median LTBI and active TB subject IFN- $\gamma$  response to culture filtrate protein was more than 10-fold higher than that of controls (Figure 1C). This suggests that most of the response to culture filtrate protein was specific to *M.tb* complex. Interestingly, we observed decreased IFN- $\gamma$  production in response to CFP in the elderly who had no history of *M.tb* infection, relative to adults (Figure 1C). Because BCG status had no effect on this observation and we do not expect significantly different prior NTM exposure between the adult and elderly controls, we believe this suggests hypo-responsiveness of the innate immune system of the elderly to the mycobacterial antigens in CFP rather than a decreased adaptive response.

We cannot rule out the possibility that *M.tb*-specific T cell function in the elderly may be impaired within the microenvironment of *M.tb* infected tissues, especially the lung. However, published data for adaptive immune responses in the lungs of aged human subjects are extremely limited. Our lab has previously shown that *ex vivo* responses to *M.tb* antigens are comparable between blood and lung among different strains of mice with different TB disease susceptibility, suggesting that our observations of no difference in *M.tb*-specific T cell function in the blood of different age groups may hold true in the lung (Beamer et al., 2008b). While previous work in an intravenous route *M.tb* infection model has shown that aged mice are more likely than young mice to die from a primary *M.tb* infection and delayed recruitment of CD4 T cells to the spleen contributes to this susceptibility, no study has demonstrated a clear defect in CD4 or CD8 T cell immunity to *M.tb* in aged mice using a pulmonary infection model (Orme, 1987; Orme et al., 1993; Wang et al., 2012).

In the case of general *in vivo* T cell immune function in elderly humans, several studies have previously shown that the impaired DTH response in the skin to recall antigens in the elderly is a general phenomenon to numerous antigens, not just PPD from *M.tb* (Castle et al., 1990; Agius et al., 2009). Agius et al have shown that in the case of *Candida albicans* antigens, the reduced skin DTH response in the elderly is due to defective activation of dermal blood vessels resulting from decreased TNF-a secretion by cutaneous macrophages (Agius et al., 2009). In this study, the intrinsic ability of peripheral blood T cells to migrate or of isolated cutaneous macrophages to secrete TNF-a upon *in vitro* stimulation were not impaired in the elderly. More recently, Vukmanovic-Stejic et al. showed that the impaired DTH to varicella zoster virus (VZV) in the elderly is associated with an antigen-nonspecific over exuberant proinflammatory response to injected saline, which was particularly associated with transient increased HLA-DR+, CD14+ and/or CD16+ mononuclear phagocytes likely recruited from the blood (Vukmanovic-Stejic et al., 2017). Contrary to the DTR to Candida albicans, early dermal blood vessel activation in response to VZV antigens was not impaired in the elderly (Agius et al., 2009; Vukmanovic-Stejic et al., 2017). Treatment with a p38 MAP kinase inhibitor significantly reduced plasma C reactive protein levels, decreased peripheral blood monocyte secretion of IL-6 and TNF-a and enhanced the VZV-specific DTR response in the skin of old subjects (Vukmanovic-Stejic et al., 2017).

Interestingly, we observed an age-dependent increase in nonclassical monocyte proportion in elderly subjects with LTBI. We hypothesize that this monocyte subset may play a role in the susceptibility of the elderly to develop active TB, possibly by impairing T cell responses at the site of infection, as in the cited studies in the skin (Vukmanovic-Stejic et al., 2017). It has been shown that in adults with pulmonary TB, the proportion of CD16+ monocytes is increased and this correlates with disease severity and TNF-a plasma levels (Balboa et al., 2011; Lastrucci et al., 2015). Moreover, this monocyte subpopulation is more permissive for *M.tb* growth, more migratory and is inhibited in the ability to stimulate T cell proliferation and cytokine production (Lastrucci et al., 2015). Therefore, we hypothesize that nonclassical monocytes in the elderly may play an excessively inflammatory role at the site of infection, be more permissive to *M.tb* growth and impair T cell responses at the site of infection. Current studies in our laboratory are ongoing to assess this hypothesis in the elderly. Our work does not preclude the possibility that the anti-mycobacterial functions of classical and intermediate monocytes are also impaired in the elderly, which has been suggested by one

study (Guerra-Laso et al., 2013). Our work has the potential to link the known widespread low-grade systemic inflammation of old age, termed "inflammaging," with the elderly's specific susceptibility to developing active TB (Franceschi and Campisi, 2014; Piergallini and Turner, 2017).

Our study is limited by sample size and available outcomes data, and thus our ability to comment on immunological correlates of susceptibility of the elderly to die from active TB is limited. We did observe a significantly increased monocyte/lymphocyte ratio in the elderly with active TB, relative to all other groups. In adults and children, an elevated monocyte/ lymphocyte ratio has prospectively been associated with risk of developing active TB and dying from disease (Naranbhai et al., 2014a, 2014b). In adults, a very low monocyte/ lymphocyte ratio has also been associated with risk of developing active TB (Naranbhai et al., 2014a). Thus, our additional observation of a decreased monocyte/lymphocyte ratio in elderly with LTBI relative to controls and adults with LTBI suggests its potential role in the risk of the elderly to develop active TB. Further research is needed to identify biomarkers that associate with poor outcomes in the elderly with TB, together with partially known risk factors for their more difficult diagnosis and treatment (Oshi et al., 2014; Velayutham et al., 2014).

Our study is one of the first in decades to study immune function in the elderly with LTBI and active TB. Importantly, our study also included a *M.tb*-uninfected cohort to account for biological effects due solely to age. With regard to *M.tb* specific T cell frequency and production of cytokines important for *M.tb* control, our data show no evidence for *M.tb*-specific T cell dysfunction in the elderly with *M.tb* infection relative to adults, and our data is suggestive of an age-dependent alteration in monocyte proportion and phenotype that may correlate with susceptibility to active TB. While we do not provide a quantification due to our low sample size, our data are also consistent with the conclusion that the QFT and T-SPOT.TB tests provide concordant results in the elderly with LTBI or active TB. Although the sensitivities of these tests have been previously compared for diagnosing active TB in the elderly (Bae et al., 2016; Du et al., 2018), no published study has directly compared these tests in the elderly population with LTBI.

Our study has several limitations. The principle limitation is our low sample size in our elderly LTBI and both active TB groups. Therefore, our results are suggestive but not definitive for the parameters we measured. Together with this limitation, it was also difficult to control for the higher prevalence of diabetes, cancer and immunosuppressive treatment in the elderly in study recruitment. However, our multivariate analyses showed that most differences we observed between age groups among controls were due to age, independent of these three confounders and others for which we have data. Importantly, diabetes, cancer and immunosuppressive treatment did not confound our conclusions for monocyte phenotypes in the elderly with LTBI, as only 1 subject from this group for whom we collected monocyte data had diabetes, and this subject had the lowest nonclassical monocyte proportion of the group (data not shown).

Additionally, in our small cohort, we could not distinguish between active TB developed soon after exposure to *M.tb*, termed primary TB, or disease which occurred years after initial

exposure, termed reactivation TB. Finally, our elderly LTBI and active TB subjects were on average treated longer with TB medications before study recruitment than the adults, and the tissue distribution of TB disease was different between our adult and elderly subjects. Our low sample size precludes our ability to estimate the effect of these differences in *M.tb* infected subjects for our monocyte data.

Overall, our study highlights the need for further research into the biological reasons why the elderly are more susceptible to TB, so that public health systems can be better equipped to face the present and future problem of TB in an aging global population.

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# Abbreviations

CFP	culture filtrate protein
CNS	central nervous system
COPD	chronic obstructive pulmonary disease
DTH	delayed type hypersensitivity
НСТ	hematocrit
NTM	nontuberculous mycobacteria
PBMC	peripheral blood mononuclear cell
PPD	tuberculin-purified protein derivative
QFT	QuantiFERON <sup>®</sup> TB-Gold Test
SFUs	spot forming units
SIADH	syndrome of inappropriate antidiuretic hormone secretion
TST	tuberculin skin test
VZV	varicella zoster virus
WBC	white blood cell count

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# Highlights

- Human immune function stratified by age and infection or disease from tuberculosis
- Tuberculosis-specific T cell number and cytokine production not impaired in elderly
- Alterations in monocyte proportion and phenotype with age and tuberculosis status

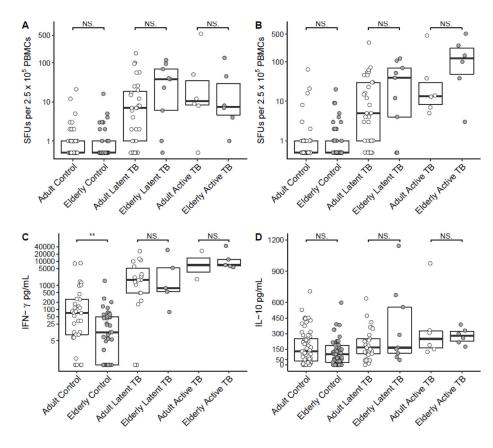
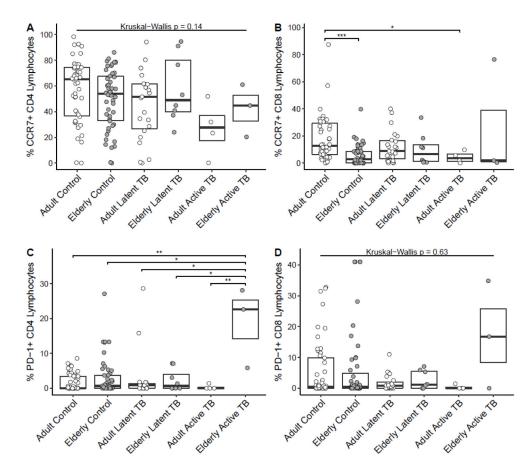
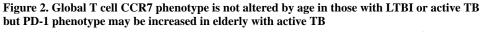


Figure 1. T cell cytokine responses to *M.tb* antigens are not altered by age in those with LTBI or active TB

Number of IFN- $\gamma$ + T cells, measured in spot forming units (SFUs), responding to ESAT-6 (A) and CFP-10 (B) from *M.tb*. IFN- $\gamma$  (C) and IL-10 (D) from culture supernatants of PBMCs stimulated with *M.tb* culture filtrate protein. Boxplots are medians with interquartile ranges. Total subjects for each panel are 157 (A), 157 (B), 100 (C) and 164 (D). \*\* p<0.01 by Mann-Whitney U test.





Percent of CD4+ (A) and CD8+ (B) lymphocytes expressing CCR7. Percent of CD4+ (C) and CD8+ (D) lymphocytes expressing PD-1. For all comparisons between age groups not displayed, there were no significant differences (p>0.05). Boxplots are medians with interquartile ranges. Total subjects for each panel are 135 (A), 134 (B), 143 (C) and 142 (D). Other Kruskal-Wallis p values were p=0.0001 (B) and p=0.03 (C). \* p<0.05, \*\* p<0.01, \*\*\*p<0.001 by Dunn's test.

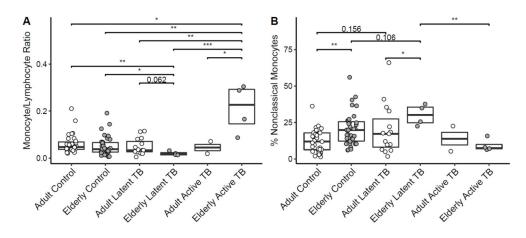


Figure 3. Monocyte/lymphocyte ratio is increased in elderly with active TB and decreased in elderly with LTBI, and old age and *M.tb* infection skew monocytes towards a nonclassical phenotype

Monocyte/lymphocyte ratio (A) and percent monocytes with nonclassical (CD14dimCD16+) phenotype (B). Boxplots are medians with interquartile ranges. Total subjects for each panel are 91 (A) and 91 (B). Kruskal-Wallis p values were p=0.002 (A) and p=0.003 (B). \* p<0.05, \*\* p<0.01, \*\*\*p<0.001 by Dunn's test, or p value otherwise displayed.

Table 1

# **Study Population Characteristics**

Due to our complex study design and available patient populations, not all covariates are balanced across all 6 study groups. p-values are from chi-squared test for categorical data and ANOVA for continuous variables.

	Adult Control	Elderly Control	Adult Latent TB	Elderly Latent TB	Adult Active TB	Elderly Active TB	All groups p-value	TB/LTBI groups p-value
n	63	53	31	6	9	L		
Sex (%)	24 (38.1)	24 (45.3)	16 (51.6)	4 (44.4)	3 (50.0)	4 (57.1)	0.818	
Age (mean (range))	34.21 (25-44)	73.23 (65–92)	31.48 (25–42)	75.56 (66–87)	33.33 (26–38)	76.29 (65–89)		
Ethnicity (%)							<0.001	0.383
Asian	6 (9.5)	2 (3.8)	17 (54.8)	4 (44.4)	1 (16.7)	5 (71.4)		
Black	10 (15.9)	7 (13.2)	9 (29.0)	4 (44.4)	3 (50.0)	0 (0.0)		
Other	0 (0.0)	2 (3.8)	3 (9.7)	1 (11.1)	2 (33.3)	1 (14.3)		
White	47 (74.6)	42 (79.2)	2 (6.5)	0 (0.0)	0 (0.0)	1 (14.3)		
Previous BCG Vaccination (%)							0.078	0.424
No	30 (47.6)	15 (28.3)	12 (38.7)	5 (55.6)	1 (16.7)	1 (14.3)		
Unknown	13 (20.6)	14 (26.4)	13 (41.9)	3 (33.3)	2 (33.3)	4 (57.1)		
Yes	20 (31.7)	24 (45.3)	6 (19.4)	1 (11.1)	3 (50.0)	2 (28.6)		
Tobacco Use (%)							0.002	0.178
No	36 (57.1)	23 (43.4)	15 (48.4)	4 (44.4)	4 (66.7)	2 (28.6)		
Previous	3 (4.8)	13 (24.5)	1 (3.2)	2 (22.2)	2 (33.3)	2 (28.6)		
Unknown	15 (23.8)	6 (11.3)	1 (3.2)	0 (0.0)	0 (0.0)	1 (14.3)		
Yes	9 (14.3)	11 (20.8)	14 (45.2)	3 (33.3)	0 (0.0)	2 (28.6)		
Alcohol Use (%)							<0.001	0.054
No	18 (28.6)	28 (52.8)	21 (67.7)	5 (55.6)	6 (100.0)	3 (42.9)		
Previous	0 (0.0)	0 (0.0)	0 (0.0)	2 (22.2)	0 (0.0)	0 (0.0)		
Unknown	17 (27.0)	6 (11.3)	4 (12.9)	0 (0.0)	0 (0.0)	2 (28.6)		
Yes	28 (44.4)	19 (35.8)	6 (19.4)	2 (22.2)	0 (0.0)	2 (28.6)		
On TB/LTBI treatment			29 (93.5)	9 (100.0)	6 (100.0)	7 (100.0)		0.688

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	Adult Control	Elderly Control	Adult Latent TB	Elderly Latent TB	Adult Active TB	Elderly Active TB	All groups p-value	TB/LTBI groups p-value
Enrolled within 30 days of TB/LTBI treatment start (%)			29 (93.5)	4 (44.4)	6 (100.0)	4 (57.1)		0.002
Diabetes Mellitus (%)	2 (3.2)	14 (26.4)	0 (0.0)	3 (33.3)	1 (16.7)	1 (14.3)	<0.001	0.020
WBC (mean (sd))	7.25 (2.58)	7.71 (2.90)	7.35 (2.68)	7.60 (2.92)	6.88 (0.69)	7.47 (2.34)	0.964	0.960
HCT (mean (sd))	40.17 (5.18)	39.61 (9.69)	43.46 (4.15)	37.11 (4.00)	38.07 (7.60)	38.11 (7.42)	0.125	0.005
Albumin (mean (sd))	4.10 (0.75)	4.03 (0.35)	4.47 (0.29)	3.76 (0.71)	4.10 (0.91)	3.65 (0.61)	0.002	<0.001
On Immunosuppressive Therapy (%)	2 (3.2)	16 (30.2)	1 (3.2)	0 (0.0)	1 (16.7)	1 (14.3)	<0.001	0.359
Positive HIV Status (%)	1 (1.6)	1 (1.9)	0 (0.0)	0 (0.0)	1 (16.7)	0 (0.0)	0.131	
Hepatitis B Infection (%)	0 (0.0)	0 (0.0)	3 (9.7)	3 (33.3)	0 (0.0)	0 (0.0)	<0.001	0.108
History of Cancer (%)	1 (1.6)	12 (22.6)	0(0.0)	0(0.0)	0 (0.0)	1 (14.3)	<0.001	
Has Other Comorbidity $\stackrel{i}{\tau}(\%)$	9 (14.3)	15 (28.3)	3 (9.7)	1 (11.1)	2 (33.3)	3 (42.9)	0.102	0.119

<sup>7</sup>Other recorded comorbidities included asthma, rheumatoid arthritis, chronic kidney disease, COPD, bronchiectasis, frequent pneumonias, emphysema, psoriasis (1 subject), depressive disorder (1), narcotic painkiller use (1), thoracic aneurysm (1), microcytic anemia-thalassemia (1), Bechet's syndrome (1), CNS Toxoplasmosis (1 Adult Active TB), SIADH (1), gastrointestinal bypass surgery (1) and liver transplant (1).

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# Table 2

# Active TB Subject Characteristics

p-value is from chi-squared test.

	Adult Active TB	Elderly Active TB	p-value
n	6	7	
Culture Confirmed <i>M.tb</i> (%)	4 (66.7)	6 (85.7)	0.879
Primary Body Sites Affected (%)			
Pulmonary	1 (16.7)	5 (71.4)	
Pleural	1 (16.7)	2 (28.6)	
Pulmonary and Musculoskeletal	1 (16.7)	0 (0.0)	
Musculoskeletal	1 (16.7)	0 (0.0)	
CNS TB and Pulmonary NTM	1 (16.7)	0 (0.0)	
Uveitis	1 (16.7)	0 (0.0)	

# Table 3

Multivariate linear models of nonclassical monocyte proportion in adult (n=31) and elderly (n=35) controls based on clinical covariates.

Independent Variable	Sum of Squares	F Value	P Value
Model 1. Dependent variable: Nonclassical monocyte proportion			
Age Group	1127.2	12.79	0.0007
Diabetes	855.5	9.71	0.0028
History of Cancer	69.2	0.79	0.38
On Immunosuppressive Therapy	66.0	0.75	0.39
Has Other Comorbidity	19.9	0.23	0.64
Residuals	5287.6		
Model 2. Dependent variable: Nonclassical monocyte proportion			
Age Group	1127.2	13.58	0.0006
Diabetes	855.5	10.31	0.0024
Ethnicity	926.6	2.79	0.037
Alcohol Use	315.7	1.90	0.16
History of Cancer	69.2	0.83	0.37
On Immunosuppressive Therapy	66.0	0.80	0.38
Has Other Comorbidity	19.9	0.24	0.63
HIV Status	17.6	0.21	0.65
Sex	13.1	0.16	0.69
Previous BCG Vaccination	38.1	0.23	0.80
Tobacco Use	76.6	0.31	0.82
Residuals	3900.0		