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### Early T-cell responses in tuberculosis immunity

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#### Summary

Tuberculosis (TB) has plagued mankind for millennia yet is classified as an emerging infectious disease, because its prevalence in the human population continues to increase. Immunity to TB depends critically on the generation of effective  $CD4^+$  T-cell responses. Sterile immunity has not been achieved through vaccination, although early T-cell responses are effective in controlling steady-state infection in the lungs. Although such early T-cell responses are clearly protective, the initiation of the *Mycobacterium tuberculosis* (*Mtb*) T-cell response occurs much later than is the case following other aerogenic infections. This fact suggests that there is a critical period, before the activation of the T-cell response, in which *Mtb* is able to establish infection. An understanding of the factors that regulate early T-cell activation should, therefore, lead to better control of the disease. This review discusses recent work that has investigated the early development of T-cell immunity following *Mtb* infection in the mouse.

#### Keywords

T cells; mycobacteria; immunity

#### Introduction

Although tuberculosis (TB) has afflicted the human population for millennia, it is nonetheless considered an emerging infectious disease, in part because disease incidence in human populations is increasing and because new forms of the pathogen pose additional threats to human health. Over two billion people are currently infected with *Mycobacterium tuberculosis* (*Mtb*), and many will experience active TB in their lifetimes (1). It is estimated that three million people will die each year from infection, making TB one of the major infectious diseases worldwide (1). TB infection rates are rising globally, particularly in Third World countries, where the combined factors of human immunodeficiency virus (HIV) infection, political instability, and breakdown of government infrastructures have contributed to declines in public health services. The problem has been further exacerbated by the increasing prevalence of multiple drug-resistant and extensively drug-resistant TB strains (2). Owing to the increasing mobility of the human population, these problems, in turn, increase the likelihood that the disease will become re-established in developed countries, where it has not been a serious problem since the onset of effective antibiotic therapies.

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There is an urgent need for more effective controls of TB worldwide. Unfortunately, currently available vaccines have only limited efficacies (3–5). Chemotherapeutic regimens are effective against drug-susceptible TB but are typically long term, leading to patient non-compliance. Although a number of promising new vaccines are under development (3, 5), real progress will require more detailed knowledge of the host immune responses. Specific areas of that require elucidation include the exact roles of various immune cell subsets, the identities of the antigens recognized by immune effector cells, the effector mechanisms involved, the mechanisms of pathogen avoidance or subversion of immunity, and the regulation of the innate and adaptive immune responses.

Only a small percentage of humans known to have been exposed to *Mtb* will develop clinical manifestations and/or symptoms of the disease (6). This peculiarity suggests that the majority of individuals exhibit innate immunity sufficient to control naturally encountered infections. It is presumed that those who show immunological evidence of exposure, in the form of skin test reactivity, harbor the bacilli in a latent form yet are able to maintain active control of the bacteria. Similarly, only a small proportion of those who harbor latent infection will develop active disease (6). Although it is clear that immunosuppression caused by HIV or by immunosuppressive or immunotherapeutic drugs is a major factor in TB recrudescence, the underlying causes of susceptibility to active disease are not known; indeed, the contributions of host and/or bacterial genetics and host–bacterial cell interactions are not well understood.

Studies of the human cellular and humoral responses during infection and disease have been undertaken using peripheral blood and cells from the lung. A substantial humoral response to *Mtb* occurs, although the significance of this response, in terms of control of the disease, is unknown (7). The cellular response in both the blood and lungs is largely inflammatory in nature, and antigen-specific cells are capable of producing macrophage-activating cytokines (8). There is also some evidence of regulatory activity [i.e. interleukin-10 (IL-10) production (9)], although the relative contributions of inflammatory and regulatory activities to the development of disease are as yet undefined. While the nature of the cellular response in exposed and diseased individuals has been characterized, the very earliest events in T-cell activation have not been studied in depth either in humans or in animal models. Because it is likely that these early immunological events strongly affect long-term cellular responses and are therefore likely to modulate disease progression, the study of these early events form the basis of the present review.

Such a characterization of early immunity will be particularly valuable for understanding events subsequent to known exposure. Cellular responses are slow to develop in humans, i.e. skin tests are not useful before 5–6 weeks after exposure (10). If we can develop tools to examine early events in humans and/or to identify early cellular events in animal models, we will be able to better understand how the cellular response is induced in humans upon exposure. One area of intense interest in this regard is the response to vaccination with live-attenuated *M. bovis* Bacille Calmette-Guérin (BCG). While this vaccine is at best only modestly effective against adult pulmonary disease, it is administrated to neonates around the world and protects against early childhood disease (11). Recent studies in South Africa demonstrate that a multi-faceted cellular response to this vaccine occurs within 10 weeks post-vaccination (12). It remains to be seen, however, whether the nature of the initial response to vaccination bears any relation to the ability of the vaccine to protect against disease.

In addition to extensive and careful studies of vaccinated infants, a range of novel approaches are being applied to the investigation of TB immunity in adults. One recent advance is the development of multi-parameter flow cytometry, which has highlighted a

heretofore unappreciated complexity of the cellular response to mycobacterial exposure (12). Another advance is the availability of major histocompatibility complex (MHC) tetramer reagents, which allow direct measurements of antigen-specific T-cell numbers in infected individuals (13, 14). Experimental approached involving animal models serve to identify mechanisms and pathways that will help to focus the work performed in human studies. A range of animal models, including mice, guinea pigs, zebrafish, and primates, have generated recent advances (15–18), and the exciting development of *in vivo*-imaging technologies in both the zebrafish and the mouse models has provided novel three-dimensional insights into cellular dynamics during mycobacterial infections (16, 19).

In this review, we focus on recent work using the highly tractable mouse model to study an issue that has long remained an enigma. As discussed above, the very earliest events in the stimulation of the immune response to aerogenically delivered *Mtb* have not been delineated. Currently, only the mouse model allows this question to be addressed experimentally. It is hoped that the recent progress made with this model will facilitate examination of the cellular response in humans, thereby leading to better insights into both disease development and immunity.

#### The mouse as an experimental model

Features of *Mtb* infection in the mouse have been studied in depth, as summarized in several recent reviews (20, 21). Inbred mouse strains exhibit a wide range of susceptibility to *Mtb* infections (22, 23) and have been used to investigate genetic contributions to disease susceptibilities (24–26). BALB/c and C57BL/6 mice have been commonly used for studies of immunity, even though these strains are relatively resistant to intravenous and aerosol *Mtb* infection (23). The C57BL/6 strain offers a major advantage for TB research, because many immunologically relevant gene-targeted mice are available on this genetic background. A general concern with regard to the mouse model, however, has been the lack of standardization among different laboratories, where variability in such key parameters as bacterial strain, inoculation route, and inoculum dose may be responsible for disparate findings. Many investigators have used intravenous and intratracheal inoculations, although it is now widely accepted that low-dose aerosol inoculation provides the most realistic model for normal human infection (27–30). Via this route, low-dose inoculations are typically designed to deliver 75–100 bacterial CFU to the lower respiratory tract (27, 29, 31, 32).

#### Innate immunity

Once delivered to the lungs, the bacilli attach to and enter host cells via any of several host receptors, including the mannose receptor, the dendritic-cell-specific intercellular adhesion molecule-3 (ICAM-3)-grabbing non-integrin (DC-SIGN), surfactant protein A, mannosebinding lectin, and the class A scavenger receptor (33). The significance of this exploitation of multiple receptors and the relevance to pathogenicity are not yet resolved, although it has been suggested that differential receptor utilization by the bacterium may affect disease susceptibility (34). However, the ability to interact with a wide number of receptors on various cell types may also be reflective of an evolutionary strategy, which has enabled *Mtb* to infect many differing cell types. Alveolar macrophages are generally considered to be the first cells infected after aerosol infection (35), and depletion of activated macrophages can lead to a dramatically altered disease progression (36). These cells presumably provide a first line of host defense, for example, they express antimicrobial peptides, i.e. cathelicidin, that may function in the early immune response (37).

As in most other types of infections, many components of the innate response are activated in *Mtb*-infected host cells. A number of studies have investigated roles for Toll-like

receptors (TLRs), in particular TLR2 (a heterodimer of TLRs 1 and 6), TLR9, and TLR4 (38–43, 44). Recent work has suggested that interaction of the bacillus with TLR2 is due in part to effects of secreted ESAT-6, via its direct interaction with TLR2 on macrophages (45). A number of chemokines are also produced in the lungs and other tissues during infection and contribute to innate immune responses (46, 47). Although there is ample evidence that Mtb can trigger innate immunity, this triggering does not occur until several days after infection, at about the time when T-cell responses are elicited (48). Thus, although *Mtb* is clearly capable of activating the innate response, it does not do so for a prolonged period following aerosol infection; this delay likely provides a critical window of opportunity in which *Mtb* can establish infection in the lungs. The delay in the activation of the innate response in the lung may be a result of a number of factors, including low pathogen exposure (i.e. 50-200 bacteria in experimental models), naturally low inflammatory responses in the lung, or active inhibition of innate immunity by Mtb. There is no clear evidence to promote any one of these ideas over the other at present, in part because the lack of tools that are sufficiently sensitive to detect early innate responses following lowdose aerosol infection.

In addition to alveolar macrophages, dendritic cells (DCs) are likely to become infected upon aerosol exposure (34, 49, 50), and these cells probably play a major role in triggering T-cell responses (51). Some investigators have proposed that the *Mtb* infection compromises DC function (52), perhaps providing a means of immune evasion. After aerosol exposure, bacteria multiply in the lungs but disseminate to lymphoid tissues and other organs. How these early growth and dissemination phases impact the generation of the T-cell response was not well studied, before some significant recent advances described below.

#### Cellular immunity

In the mouse, *Mtb* grows in an apparently uncontrolled fashion for about the first 21 days (53), at which time further increase in steady-state bacterial numbers come under the control of the immune system (31, 54). A stationary phase of bacterial colonization ensues; immunity during this period is maintained by T cells until the animal succumbs to infection, an event which in C57BL/6 mice occurs between 200 and 250 days after infection (31, 54). The precise cause of death is not known, although it could be a consequence of a loss of cellular immunity or a result of chronic T-cell-mediated pathology.

In the mouse model, the roles of particular immune cell subsets, cytokines, and chemokines during host adaptive immunity have been studied in depth. It is now well established that CD4<sup>+</sup> T cells are instrumental in the control of TB infections in both humans and mice (21, 55–58). Chief among the functions of CD4<sup>+</sup> T cells is the production of interferon (IFN), which is required for resistance to fatal infection, given that both class II MHC- and IFN deficient mice rapidly succumb to infection (59, 60–63). An important function of IFN is the induction of nitric oxide synthesis in macrophages; IFN in turn controls bacterial growth (63–66). Although  $CD4^+$  T cells are widely assumed to provide an essential source of IFN, this relationship has not been formally demonstrated. Some studies have suggested that immune control is mediated independent of IFN (60, 67). Moreover, although CD4<sup>+</sup> T cells are required for control of infection in the mouse, these immune cells are unable to eliminate the pathogen or to prevent fatal disease. T-helper 1 (Th1) responses are nevertheless a major component of protective immunity, because they clearly limit bacterial expansion. CD4<sup>+</sup> T cells are well known to be required for immunity to Mtb, and recruitment of T cells to the lung is necessary for containment of *Mtb* within granulomas (68).

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impervious cell wall, is resistant to otherwise damaging anti-microbial compounds produced by the host cell (i.e. reactive nitrogen and oxygen intermediates). Alternatively, the bacterium could actively subvert or modify effector functions of macrophages, DCs, T cells, or other host cells (69–71). It is also possible that the inflammatory consequences of infection produce an internal or external environment that is not conducive to bacterialkilling activity by the immune response. Currently, there are insufficient data to support any one of the above hypotheses over the others.

While CD4<sup>+</sup> T cells are known to be essential, other T cells and lymphocytes also play important roles in host defense, even when their functions have been shown to be redundant (21, 72). A number of investigators have investigated the role of CD8<sup>+</sup> T cells. In contrast to the dramatic impact of the absence of CD8 coreceptor-expressing cells on the development of disease following a high-dose intravenous infection (73), the absence of CD8<sup>+</sup> T cells in the low-dose aerosol model had little impact disease progression, until after as long as 200 days (54, 74–76). Moreover, although transfer of Lyt-2<sup>+</sup> (CD8<sup>+</sup>) cells was not particularly effective against low-dose aerosol infection, CD8<sup>+</sup> T-cell transfer was more effective than were CD4<sup>+</sup> T cells against high-dose infection (77). Thus, while CD4<sup>+</sup> T cells are able to control low-dose infection, CD8<sup>+</sup> T cells possess a unique activity that makes them essential when a high-dose infection occurs.

A difficulty in resolving the exact role of class I MHC-restricted CD8<sup>+</sup> T cells is in part due to the fact that a variety of strains of gene-targeted mice have been used to assess infection in the absence of CD8<sup>+</sup> T cells. Moreover, many gene-targeted strains exhibit incomplete elimination of the targeted cell populations, and/or they exhibit pleiotropic effects of the targeted mutations. For example, 2-microglobulin (2m)-deficient mice lack both MHC class Ia- and MHC class Ib-restricted CD8<sup>+</sup> T cells, and CD8-deficent mice contain class I MHC-restricted, CD8<sup>-</sup> T cells. Studies in the 2m-deficient mice are further complicated by an iron overload defect that renders this strain more susceptible to *Mtb* infection (78). Furthermore, the absence of 2m results in an early but limited reduction in the ability of the mice to control bacterial growth (75). However, no difference in disease development was noted in mice lacking another 2m-dependent class Ib molecule, CD1d (75, 79). Urdahl et al. (80) addressed roles for class Ia- and class Ib-restricted CD8<sup>+</sup> T cell; using class I MHC K<sup>b</sup>D<sup>b</sup>-deficient mice, they demonstrated that class Ib-restricted CD8<sup>+</sup> T cells, although present in infected lungs, were unable to fully compensate for the lack of class Ia-restricted T cells. These data revealed a non-redundant role for class Ia-restricted CD8<sup>+</sup> T cells. Like CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells have been shown to function by producing IFN and by targeting infected cells for cytolysis (81–84). More recently, it has been demonstrated that CD8<sup>+</sup> T cells recognized *Mtb* antigens generated by cross priming *in vitro* (85).

#### Mtb antigens recognized by CD4+ T cells

Early studies of T-cell immunity focused on polyclonal T-cell populations that were elicited by *Mtb* infection and re-stimulated *in vitro* with either live bacteria or crude proteins preparations. In these types of analyses, detection of antigen-specific cells was not possible (81, 86). *Ex vivo* stimulation of cells from humans, guinea pigs, and mice, using purified proteins demonstrated that immunodominant responses were associated with *Mtb* proteins, and the responses were not limited to a single *Mtb* protein (87, 88). Identification of the *Mtb* antigens and epitopes recognized by both mouse and humans (89–92) has been facilitated by the search for novel T-cell antigens relevant to vaccine development. This has led to the discovery of a wide array of *Mtb* antigens that activate CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, nonclassical class I-restricted CD8<sup>+</sup> T cells and T cells (92). These T-cell ligands include not only conventional peptide antigens that bind to classical MHC class I and class II molecules but also a number of abundant mycobacterial lipids that are presented by the non-classical class I molecule CD1 (93, 94).

Early studies suggested that immunity to *Mtb* is generated by viable but not killed bacteria (95). One explanation for these findings is that killed bacteria do not secrete *Mtb* antigens *in vivo*. Other infection models have shown antigen secretion to be an important requirement for the generation of T-cell ligands (96). Indeed, many *Mtb* proteins are released during bacterial culture (92), suggesting that such proteins are also secreted while the bacteria are residing within host cells *in vivo*. These culture filtrate proteins (CFPs) were found to be mixtures of secreted proteins and outer cell wall proteins that have been shed or released from dead bacteria (97), and considerable effort has been expended on their characterization. Such work identified many *Mtb* antigens among the CFPs that could elicit responses from T cells in both mice (87, 98) and humans (99). Importantly, CFPs have been shown to be the first antigens recognized during infection (100, 101), and lymphocyte responses are hypothesized to be directed largely against these proteins (90, 100, 102). Of considerable importance are studies in which immunization with CFPs isolated from short-term cultures elicited at least partial protection to *Mtb* infection in experimental mouse models (87, 101).

Many CFPs have now been shown to be recognized by T cells (87). One of the major CFP antigens is Ag85B (103, 104), a member of the p85 complex encoded by *fbpB* (105). The concentration of Ag85B within CFPs was 100-fold higher than concentrations in bacterial sonicates (106). Activated CD4<sup>+</sup> T cells have been shown to respond to a particular Ag85B peptide epitope in C57BL/6 mice (residues 240–254) (107). Another major CFP antigen that has been characterized is the early secreted antigenic target-6 (ESAT-6). ESAT-6 is a low molecular mass (6 kDa) protein found in abundance in *Mtb* culture filtrates (91). Because ESAT-6 is a major target of both CD4<sup>+</sup> and CD8<sup>+</sup> T cells in mice and humans, it is a major vaccine candidate (108). Immunization of mice with an ESAT-6 peptide induced protection equivalent to that obtained by the use of BCG (109). Modified BCG strains are now available that express ESAT-6; re-introduction of the genetic region of difference RD1, which includes ESAT-6, into BCG restored pathogen virulence [the ESAT-6 gene is deleted in the attenuated BCG strain (111)] (110).

The identification of new *Mtb* T-cell epitopes is proceeding at a rapid pace, and the new antigens will be of use for studies of cellular immunity. A recent study has cataloged all currently known *Mtb* T- and B-cell epitopes identified in a wide range of mammalian species (112). Given that only a small portion of the *Mtb* proteome has been studied, many additional T-cell epitopes in different host species will undoubtedly be identified. Nevertheless, the epitope database, along with other studies of *Mtb* T-cell epitopes (72, 89), will be an important tool for both researchers and clinicians.

#### Expression of Mtb antigens in vivo

Given the identification of multiple *Mtb* T-cell antigens, it will be important to ascertain when and where these antigens are produced during infection. This information is critical for understanding the genesis and maintenance of protective immunity, especially because antigen expression likely varies temporally and spatially during infection. One study investigated the expression of ESAT-6 and Ag85B following virulent *Mtb* (H37Rv) infection (113). mRNAs for both antigens were expressed during acute and chronic phases of infection, although the magnitude and kinetics of antigen mRNA expression differed. Both mRNAs were expressed early following infection, but ESAT-6 mRNA expression persisted at 10-fold higher levels through day 20 post-infection. Expression of both antigens declined after day 20; based on the ratio of transcripts to *Mtb* CFUs, only a few of the

bacteria expressed these antigens *in vivo*. These data also suggest that the bacilli enter into a quiescent, non-dividing state *in vivo*, once their expansion has been controlled by the cellular immune response. These data raise an important question: how are protective long-term effector T-cell responses, which likely require a stable source of antigen, maintained during chronic and/or latent infections. Although the study by Rogerson *et al.* (113) showed evidence of major changes in mRNA expression that could contribute to differing T-cell responses during acute and chronic infection, the work needs to be extended through study of additional *Mtb* antigens.

#### Analyses of antigen-specific T cells

The identification of the ESAT-6 peptide epitope recognized in C57BL/6 mice made available a powerful tool that allowed us to examine when and where IA<sup>b/</sup>ESAT-6-specific CD4<sup>+</sup> T cells are located during acute and chronic phases of infection (114). In our studies, we used enzyme-linked immunospot (ELISpot) analyses to enumerate IA<sup>b</sup>/ESAT-6<sub>1 - 20</sub>specific CD4<sup>+</sup> T cells in vivo. (Note that because the secretion of IFN is antigen dependent, this approach does not reveal whether IFN is actually expressed *in vivo* but rather measures the capacity of the cells to produce cytokines following antigen stimulation). We observed that  $IA^{b}/ESAT-6_{1-20}$ -specific T cells were detectable no earlier than 14 days post-infection in the lungs and the mediastinal lymph node (MLN). High frequencies of responder cells were not detected until day 21, and the response was maximal on day 28 post-infection in the lungs. At that time point, the frequency of antigen-specific T cells was approximately 1.5% of total lung CD4<sup>+</sup> T cells, corresponding to  $1-5 \times 10^5$  cells in the lungs. Thereafter, the frequency of responding cells stabilized in the lungs at approximately 1%, a frequency maintained at least through day 160 post-infection. Similar findings were obtained in the MLN and spleen, except that the frequencies of T cells in these latter tissues were lower. These data revealed that the antigen-specific effector CD4<sup>+</sup> T-cell response did not become detectable until relatively late following infection, that the CD4<sup>+</sup> T cells underwent major expansion, and that chronic infection was associated with a persistent T-cell presence in the lungs. The antigen-specific cells identified in this study exhibited characteristics of fully activated effector cells: high expression of CD44, low expression of CD62L, and the ability to secrete IFN upon contact with antigen (114). While we concluded that cellular immune responses must be present within the lung, the extent to which the lung CD4<sup>+</sup> T cells play a role in maintaining control of bacterial burden during the chronic phase is unclear.

Several other investigators drew similar conclusions regarding the generation of antigenspecific effector  $CD4^+$  T cells. Chackerian *et al.* (115) observed IFN production by a spleen cell population in response to a sonicate of *Mtb* following aerosol infection, as early as day 11 post-infection in C57BL/6 mice. The particularly early response in that study could have resulted from the use of the complex antigen preparation and/or the production of IFN by non-T lymphocytes. Lazarevic *et al.* (116) reported that IA<sup>b</sup>/ESAT-6-specific T cells undergo dynamic changes during acute and chronic infection. The above studies have made important contributions to the study of early T-cell immunity, by revealing that the onset of the T-cell response during *Mtb* infection is delayed, compared with infections mediated by other pulmonary pathogens such as the influenza virus (117).

#### Antigen-specific CD8+ T-cell responses

Several investigators have examined antigen-specific CD8<sup>+</sup> T-cell responses. In one study, CD8<sup>+</sup> T-cell responses were shown to be initiated in the lung-draining lymph node (LN) within 11 days after infection (118). The T-cell response was also interpreted to be initiated only after the dissemination of *Mtb* to the MLN, given that bacterial infection was first detected on day 9 post-infection. Thus, the authors concluded that bacterial dissemination

precedes the development of the T-cell response. However, a caveat relates to the sensitivity of the IFN enzyme-linked immunosorbent assay (ELISA) used to detect T-cell responses; possibly the T cells had undergone activation before day 9 post-infection, but IFN was present at levels below the limits of detection.

Other studies have examined CD8<sup>+</sup> T-cell responses to defined class I MHC peptides, including K<sup>d</sup>/TB10.3/10.4<sub>20-28</sub> (119) and K<sup>k</sup>/CFP<sub>32-39</sub> (83). The K<sup>k</sup>/CFP<sub>32-39</sub> specific cytolytic CD8<sup>+</sup> T cells were first detected 2-3 weeks post-infection and persisted in the lungs for as long as 260 days post-infection (83). These studies of antigen-specific effector CD8<sup>+</sup> T cells were extended after the development of class I MHC tetramers that recognize K<sup>k</sup>/CFP<sub>32-39</sub>- and K<sup>d</sup>/TB10.3.4/10.4<sub>20-28</sub>-specific CD8<sup>+</sup> T cells (83). Remarkably, K<sup>d</sup>/ TB10.3.4/10.4<sub>20 - 28</sub>-specific T cells were found at frequencies as high as 30% in the lungs of chronically infected BALB/c mice, demonstrating that TB10.3 and TB10.4 are immunodominant antigens in this strain. An additional MHC class I tetramer has been described that recognizes the Mtb72F polyprotein in the context of the MHC class II protein IA<sup>b</sup> (120). In that study, the tetramer was used to identify IA<sup>b</sup>/Mtb72F<sub>92-102</sub>-specific CD8<sup>+</sup> T cells in aerosol-infected mice within 30 days and as late as 100 days post-infection. The  $IA^{b}/Mtb72F_{92-102}$ -specific CD8<sup>+</sup> T cells were present at frequencies as high as 7.5% in low-dose-infected mice, indicating that Mtb72 is also an immunodominant antigen and that the antigen is expressed during chronic infection. The availability of mouse MHC class I tetramer reagents is thus a significant advance and will be particularly useful for future analysis of the antigen-specific lymphocyte responses during the adaptive immune response to Mth.

#### T-cell responses are delayed in *Mtb* infections

The relatively late initiation of the CD4<sup>+</sup> and CD8<sup>+</sup> T-cell responses to aerosol Mtb infection contrast with the timing of the responses to several acute viral infections (121, 122) as well as with the timing of responses to other intracellular bacterial infections (123–125). In these other infection models, antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T-cell responses were typically detected much earlier than are detected during *Mtb* infection, often as early as 5 days post-infection. Although they occur late following aerosol inoculation, Mtb-specific Tcell responses are not inherently delayed, because experimental intravenous inoculation can generate robust T-cell activation within 1–3 days post-infection (51, 126, 127). Our recent studies, in which  $2 \times 10^5$  bacteria were administered intravenously, revealed that priming occurred at as early as 18 h (W. Reiley, G. Winslow and D. Woodland, unpublished data). Such early T-cell priming was observed by other investigators, who examined T-cell responses to an ovalbumin epitope expressed by a recombinant BCG (128). In those studies, transgenic OVA-specific CD4<sup>+</sup> T cells initiated cell division within 5 days of infection, supporting the idea that early CD4<sup>+</sup> T-cell responses can occur early following BCG infection, if sufficient antigen is available. A limitation of the above studies of early antigenspecific T cells responses is that none had been able to identify the exact time, after infection, when T-cell responses are initiated. The inability has been in part due to the lack of tools that would allow the direct detection of T-cell responses in vivo. For example, studies have been limited by reliance on indirect assays of T-cell activation (i.e. cytokine secretion); these effector responses require 3-4 days to develop after the T cells first encounter antigen.

Why, then, are T-cell responses to *Mtb* detected relatively late after infection? This question is an important one, because a delay in the generation of cellular immunity is likely a critical factor in allowing the bacteria to establish persistent infection. A simple explanation is that the apparent delay in T-cell activation is due to the lack of MHC/peptide complexes able to prime T cells. Although the population doubling times of *Mtb* strains vary, the doubling

time within the lungs in the first 10 days following intravenous infection is approximately 36 h for strain H37Rv (53). Aerosol inoculation with 200 CFUs of H37Rv would in this case be expected to yield approximately 2000 live bacteria within the lungs within the first 5 days and 20 000 bacteria in the first 10 days post-infection. Although T cells require as few as 10-100 MHC/peptide complexes to be presented on an antigen-presenting cell (APC) before they can undergo priming (129), it is not yet possible to correlate the number of bacteria to the antigen loads on APCs; tools to quantitate MHC/peptide complexes on the surface of APCs are not available for *Mtb* antigens. Nevertheless, one study reported that effector Tcell responses in the spleen did not occur when fewer than 250 CFUs were detected (118), suggesting that the bacteria were unable to generate sufficient MHC/peptide complexes to trigger T cells. A caveat is that it is not known whether the T cells that initiate the response need to be activated by DCs that are infected by bacteria or whether DCs can acquire antigens passively, e.g. by acquiring soluble antigens or antigens acquired from apoptotic cells (85). Nevertheless, antigen availability within secondary lymphoid tissue is a likely factor that limits T-cell priming. If that is the case, it should be possible to accelerate T-cell activation by increasing the size of the experimental inoculation. Experiments designed to test this idea are described below.

It is alternatively possible that the apparent delay in the priming of the T-cell response occurs because *Mtb* avoids immune detection in infected alveolar macrophages and is, therefore, able to replicate to high numbers. Infected alveolar macrophages are unlikely to migrate to secondary lymphoid organs to initiate T-cell priming events in the lung-draining LN. It is also possible that bacteria administered during low-dose aerosol infection are deposited in a location that does not allow host cell detection and transport to the LN by DCs that are sampling the airways. Thus, an understanding of the kinetics and location(s) of the first peptide/MHC complexes available for recognition is key to deciphering the early events in T-cell activation that follow aerosol exposure to *Mtb*.

A third explanation for the apparent delay in T-cell priming is that *Mtb* suppresses early antigen presentation in secondary LNs or influences the capacity of T cells to undergo activation. For example, cells loaded with mycobacterial antigens may not be capable of initiating T-cell activation after they have migrated from the lung to the draining LNs. DCs, which are infected by *Mtb*, are well known to be the most important APCs in initiation of T-cell responses *in vivo*; in this hypothesis, DC function is impaired following *Mtb* infection. Such impairment could arise, for example, through compromised DC trafficking from the lungs to the draining LNs or through a compromised ability of DCs to provide proper signals for T-cell activation. Evidence is available to support both mechanisms. Several studies have demonstrated that antigen presentation can be suppressed in macrophages and by extension DCs, following *Mtb* infection of APCs (70, 130–132). It has also been suggested that that infected *in vitro*-generated DCs can be impaired in their ability to prime T cells (52), although other studies suggest that DCs are highly effective APCs during *Mtb* infection *in vivo* (133).

#### Early events in T-cell activation

To understand how protective T-cell responses are initiated during *Mtb* infection, it is helpful to first review what is known from other infection models regarding early T-cell activation. Precedents are available from several studies of early T-cell responses following acute viral infections (117, 121). Naive antigen-specific T cells migrate throughout the circulation in mice, and encounter with antigen typically occurs in LNs that drain infected tissues. The antigen-specific T cells form stable contacts, engage DCs, and undergo an orchestrated series of interactions with DCs (134–136). The initial contact, duration, and quality of the interaction made with antigen-laden DC cells are likely important factors that

influence T-cell responses (137, 138). Within 4–8 h, antigen-activated T cells express a number of important cell surface receptors, including CD69 and CD25 (the IL-2 receptor chain). CD69 is the earliest T-cell activation marker expressed and is first detected within 3 h of contact with antigen (139). After the expression of CD69, activated T cells become less mobile, losing their ability to migrate out of the LNs, due to loss of the expression of sphingosine-1-phosphate receptor-1 (S1P)(140). During the following 2 days, a programmed series of events occur within the activated T cell, leading to the upregulation of CD44, the downregulation of CD62L, and the onset of cell division. As activated T cells begin to proliferate, they become motile and again encounter DCs (134). CD69 expression is then lost from the cell surface, and S1P<sub>1</sub> is re-expressed, enabling migration of the T cell. In the periphery, the activated T cells again encounter antigen, most likely via interactions with macrophages and DCs, and elaborate effector activities that contribute to pathogen clearance. Thereafter, most T cells die; at the population level, this results in a contraction of the response. Some T cells, in a process governed by signals that are not well defined, develop into memory T cells that can provide long-lived protection (141).

#### Early T-cell activation during *Mtb* infection

The processes of T-cell activation, differentiation, and migration during *Mtb* infection have only begun to be studied. An important advance is the availability of tools that allow the detection of endogenous antigen-specific T cells. *Mtb*-specific class I MHC tetramers, for example, described above, offer significant promise, especially if incorporated into methods that enhance the detection of very low numbers of endogenous antigen-specific cells (142).

An alternative approach to the study of T-cell activation involves the use of T cells from T-cell receptor transgenic mice. T cells from these mice form a homogenous pool of identical antigen-specific cells and have long been used to study naive T-cell responses *in vivo*. One well-described experimental technique involves the seeding of mice with naive transgenic T cells, at cell numbers sufficient for detection by flow cytometry (143, 144). Although a number of concerns have been raised recently regarding the effects of introduction of non-physiological numbers of T cells in such experimental models, the approach can be powerful, as long as possible artefacts associated with relatively high precursor cell frequencies are taken into account. Although high precursor frequencies are well known to influence effector T-cell differentiation and memory development (142, 145), transgenic T cells are valuable tools in the study of early T-cell activation *in vivo*.

An *Mtb* Ag85B-specific T-cell receptor transgenic mouse has been used by Wolf *et al.* (146) to monitor early T-cell expansion after aerosol infection. This CD4<sup>+</sup> T-cell receptor transgenic mouse (P25 TCR-Tg), which produces T cells that recognize IA<sup>b</sup>/Ag85<sub>240 – 254</sub>, was developed by Tamura *et al.* (147). In the studies of these Ag85-specific transgenic T cells, Wolf *et al.* (146) adoptively transferred  $3 \times 10^6$  IA<sup>b</sup>/Ag85<sub>240 – 254</sub>-specific CD4<sup>+</sup> T cells into naive C57BL/6 mice before aerosol *Mtb* infection. The T cells then underwent significant cell division between days 11 and 14 post-aerosol infection in the MLN, demonstrating that T-cell response to *Mtb* is initiated in this lymphoid tissue. T cells were not detected in the lungs until day 17 post-infection, and, because most of these cells had undergone many cycles of cell division, the authors concluded that those T cells were first activated in the draining LN and only thereafter migrated to the lungs. These data, in contrast to previous findings, suggested that the lungs are not a site of T-cell priming; instead, cells are activated and undergo cell division only in the MLN (148). These latter studies support the notion that recognition of antigen in the draining LNs is the critical event in the emergence of immunity to *Mtb*.

In these same studies by Wolf et al. (146), a strict correlation was observed between the number of colonizing bacteria in the mediastinal and inguinal LNs and the percentage of proliferating T cells, revealing that T-cell activation is contemporaneous with bacterial dissemination. This conclusion supports data by Chackerian et al. (118), discussed above. On the basis of these studies, it was suggested by Wolf et al. (146) that bacterial dissemination is essential for T-cell activation and that Mtb peptide/MHC antigens must be produced in the local draining LN. This proposal argues against the hypothesis that uninfected DCs acquire antigen passively in the lungs and migrate to the draining LN, where they initiate T-cell responses. An alternative explanation is that infected host cells (i.e. DCs or other APCs) traffic the bacteria to the LNs, where the migrating cells either present antigen to T cells directly or else indirectly via another APC. Although DCs were shown to migrate to the draining LNs during Mtb infection, Wolf et al. (146) argued that this process is impaired at an early stage after infection. In their interpretation, T-cell responses are delayed, not due to lack of antigen but rather to a lack of antigen-bearing DCs within the draining LNs. Such a mechanism, if correct, would be highly significant for our understanding of early T-cell immunity and protection from Mtb infection.

One prediction of the 'impaired DC' hypothesis is that impairment would occur in a fashion largely independent of changes in bacterial numbers and/or antigen load. With this in mind, Wolf *et al.* (146) addressed whether the early onset of the T-cell response was dependent on the size of the aerosol *Mtb* inoculum. They found that a reduction of the *Mtb* inoculum had no effect on the timing of T-cell activation (above a minimal inoculum threshold). In contrast, increasing the inoculum sevenfold (from 100 to 700 CFU) led to an increase in the frequency of cells that had undergone cell division on day 14. The authors argued on this basis that *Mtb* actively subverts early effector T-cell responses and in this way gains a foothold, before the development of T-cell immunity.

Although the studies described above made important contributions to our understanding of early T-cell activation following *Mtb* infection, a limitation was that T-cell activation was not directly examined. Instead, indirect readouts of T-cell activation (i.e. T-cell proliferation and cytokine secretion) were monitored. These effector T-cell responses typically are initiated only several days after T cells encounter antigen and are often accompanied by T-cell migration. Thus, in the experiments by Wolf *et al.* (146), it was necessary to infer the time of T-cell activation on the basis of the number of cell divisions undergone by the T cells.

To directly identify the earliest T-cell activation events following low-dose aerosol infection, our laboratories utilized another CD4<sup>+</sup> T-cell receptor transgenic mouse strain that produces T cells that recognize the major MHC class II IA<sup>b</sup> T-cell epitope in ESAT-6 (149). The epitope (IA<sup>b</sup>/ESAT-6<sub>1 – 20</sub>) was first identified by Brandt *et al.* (149). We generated T-cell hybridomas that recognized this epitope, characterized, and cloned the T-cell receptor and chain genes, and generated transgenic mice. The IA<sup>b</sup>/ESAT-6<sub>1-20</sub>-specific T cells recognized the antigen *in vitro* and *in vivo* during *Mtb* infection (149).

We used adoptive transfer of transgenic T cells to address exactly when CD4<sup>+</sup> T cells first become activated *in vivo*. This approach required that sufficient numbers of T cells be present in the LNs, lungs, and other organs, to allow detection and characterization of the T cells by flow cytometry. For this reason, we transferred  $0.25 \times 10^6$  and  $2 \times 10^6$  highly purified transgenic CD4<sup>+</sup> T cells to Thy-1 congenic mice and examined activation marker expression on the donor cells daily, beginning on day 8 following aerosol infection with 75 CFU of *Mtb* strain H37Rv. On the basis of CD69 upregulation, we determined that the donor T cells first encountered antigen on day 10 post-infection in the MLN. CD69 expression was never detected in transgenic T cells in the lung, suggesting that these T cells

migrated to the lungs after their initial activation in the MLN (149). Given that the timing of the earliest T-cell activation was coincident with the arrival of bacteria in the MLN, our data support the conclusions of Wolf *et al.* (146), that early T-cell priming to *Mtb* occurs exclusively in the MLN. In contrast, our direct measurement of  $IA^b/ESAT-6_{1-20}$ -specific T cells revealed that activation occurred by day 10 post-infection, 1 day earlier than had been reported for the  $IA^b/Ag85$ -specific T cells. The disparity in the findings may be due to the differences in the experimental approaches between the two studies or to differences in the kinetics of antigen presentation between Ag85B and ESAT-6.

The transgenic IA<sup>b</sup>/ESAT-6<sub>1 – 20</sub>-specific T cells used in our studies subsequently upregulated CD44 and downregulated CD62L, findings that are indicative of T-cell activation (149). These two markers were also expressed in the lungs beginning on about day 14 in both the MLN and lungs, indicating that about 4 days were required for cell division to be initiated after antigen encounter. We were unable to detect any increase above background in CD69 expression by endogenous CD4<sup>+</sup> T cells in the MLN, lungs, or spleen, on day 10–15 post-infection; the inability to detect endogenous T-cell priming validates the use of the transgenic T cells for detecting early T-cell activation (149). A schematic that summarizes the kinetics of early CD4 T-cell activation in different tissues following aerosol *Mtb* infection is shown in Fig. 1.

In contrast to the findings of Wolf *et al.* (146), we (149) observed a direct correlation between the size of the *Mtb* CFU inoculum and the timing of early activation of the transgenic T cells. Increasing the size of the aerosol inoculation approximately 15-fold (from 75 to 1200 CFU) resulted in a 2-day acceleration of T-cell activation within the lungdraining LN (149). Because the larger inoculum likely increased the available antigen in the lung, this finding suggests that antigen is a critical factor that limits the early activation of CD4<sup>+</sup>T cells. The implication of these findings is that early T-cell priming is dependent on available antigen and is not necessarily affected by inhibition of DC function and T-cell priming by the bacterium. An alternative explanation is that the higher bacterial burden induced more rapid DC migration.

On the basis of the studies discussed above (118, 146), it has been proposed that *Mtb* dissemination from the lung to the draining LN precedes and is essential for the initiation of the T-cell response. Our studies corroborate the findings of these groups, as we have observed that T-cell priming in the MLN occurs at the same time that bacteria are found in this tissue. *In vivo* depletion of DCs using CD11c-promoter diptheria toxin receptor transgenic mice delayed the spleen CD4<sup>+</sup> T-cell response and exacerbated mycobacterial infection following intravenous inoculation (51), suggesting that DC migration from the lungs to the draining LN is a critical step in T-cell priming. By comparing CFUs and the time of early T-cell activation, Wolf *et al.* proposed that T-cell activation requires that at least 1500 bacteria be present in the MLN (52). It is not known, however, if DCs or other cells are responsible for delivering the bacteria to the draining LNs or whether uninfected DCs can also acquire antigen from infected cells in the lungs before trafficking to the LN. Thus, the relationship between DC antigen delivery and bacterial dissemination is not yet resolved.

#### Temporal and spatial arrangements of lymphocytes

While flow cytometric studies provide critical information on the phenotypes, numbers, and functions of cells in different tissues, they cannot generate information on the threedimensional interactions of cells within inflamed tissues. To begin to acquire knowledge of immune responses to *Mtb in situ*, several investigators have begun to study where T cells and APCs localize within infected lungs (150–152). For example, CD4<sup>+</sup> and CD8<sup>+</sup> T cells

exhibit different spatial and temporal distributions in the lungs during infection that may contribute to or dictate cellular functions (150). We have begun to characterize the spatial localization of the transgenic IA<sup>b</sup>/ESAT-6<sub>1 – 20</sub>-specific T cells in our experimental model. Within 17 days post-infection, the transgenic cells were visualized in the lung airways in infected mice (Fig. 2). Other investigators have reported that chronic infection in both mice and humans is associated with the development of chemokine-dependent lymphoid-like structures within the lungs (153, 154). Such studies are likely to be important for understanding the interaction of T cells with APCs and other cells *in vivo*.

#### **Events following T-cell activation**

Once activated, the IA<sup>b</sup>/ESAT-6-specific T cells regulated the expression of several markers associated with T-cell differentiation, including CD62L, CD27, CD43, and CD127, consistent with the differentiation of the newly activated T cells into effector T cells (W. Reiley, G. Winslow and D. Woodland, unpublished data). The CD4<sup>+</sup> T cells exhibited a normal pace of T-cell differentiation, suggesting that CD4<sup>+</sup> T-cell responses are unabated during Mtb infection. Other investigators have reported similar findings from studies of polyclonal populations of CD4<sup>+</sup> T cells during *Mtb* infection (155). These data suggest that Mtb does not impair T-cell responses, once the T cells have been activated. However, two recent studies have reported, in contrast, that early CD4<sup>+</sup> T-cell responses in fact may be impaired, due to the activity of forkhead box protein 3 (Foxp3) and/or CD25<sup>+</sup> T-regulatory cells (156, 157). In the study by Urdahl et al. (156), Foxp3+ T-regulatory cells were detected in the lungs and LNs of *Mtb* infected mice within 21 days post-infection, although T-cell numbers increased significantly later, during chronic infection. In the study by Kursar et al. (157), naive CD4<sup>+</sup> T cells were transferred to recombination-activating gene (RAG)deficient mice, with or without putative CD25<sup>+</sup> T-regulatory cells, before *Mtb* infection. Expansion of the naive CD4<sup>+</sup> T cells was inhibited in the presence of the co-transferred CD25<sup>+</sup> T cells. In both studies, bacterial infection was higher in mice that contained putative T-regulatory cells, a fact that argues for a role of these cells in attenuating T-cell responses during both acute and chronic infection. Remarkably, transfer of CD25<sup>-</sup> CD4<sup>+</sup> T cells to RAG-deficient mice in the absence of the CD25<sup>+</sup>T-cell population resulted in a  $1-2 \log_{10}$ reduction of bacterial infection in lung and spleen. These studies together suggest that Tregulatory cells play a critical role in attenuating T-cell responses during *Mtb* infection, although clearly additional studies are warranted.

#### Chronic Mtb infection

Although our focus in the present review is on early T-cell activation, mouse models are also valuable for the study of chronic *Mtb* infection. As discussed above, bacterial infection is stabilized at about  $1 \times 10^6$  bacteria per lung after about 28 days post-infection, at least in part due to the action of CD4<sup>+</sup> T cells. During this period of chronic infection, which can last as long as 250 days, antigen-specific effector CD4<sup>+</sup> T cells are actively recruited and maintained in the lungs (114, 158). Our study showed that the steady-state CD4<sup>+</sup> T-cell population under went continual turnover, during which the entire population was replaced within 7 days; the rate of turnover declined once control of infection was established (114). These data suggest that CD4<sup>+</sup>T cells are either maintained by proliferation, within the lungs or the lymphoid organs, or else are continually recruited from naive thymus-derived T cells. The former possibility was suggested by our finding that the lung IA<sup>b</sup>/ESAT-6-specific Tcell population was maintained in thymectomized mice, although a caveat is that conventional thymectomy may have not eliminated all thymic tissue in that study (159). The data nevertheless reveal that sufficient IA<sup>b</sup>/ESAT-6<sub>1-20</sub> was present throughout chronic infection to maintain long-term effector T-cell responses. This observation does not rule out a contribution from newly generated T cells to immunity during chronic infection, however.

In normal mice and humans, naive T cells are produced lifelong by the thymus, albeit at agedependent rates. These newly formed T cells may be recruited to chronically *Mtb*-infected tissues and/or granulomas and may contribute to the control of infection, although this idea has not been formally investigated. We have begun to model a possible contribution of naive CD4<sup>+</sup> T cells during chronic infection, by transferring IA<sup>b</sup>/ESAT-6<sub>1 – 20</sub>-specific transgenic T cells to mice during various stages of infection.

#### Memory T cells and vaccination

Investigators have used a range of different immunization strategies to address vaccine efficacy in mice; in a number of cases, a degree of protection has been achieved (typically 1  $\log_{10}$  reduction in bacterial load) (20, 127, 160). Such studies have provided an experimental rationale for human vaccine trials (3), although researchers have only begun to identify human memory phenotype T or B cells generated by vaccination (161). In the mouse model, administration of defined antigens and adjuvants has been shown to result in the early recruitment of CD4<sup>+</sup> T cells to the lungs upon challenge, resulting in better control of infection (152, 158, 162). Those latter studies have suggested that effector T cells can be elicited earlier than in unimmunized mice following *Mtb* inoculation and that these earlier responses are critical in establishing immune control of infection. Early T-cell responses act to limit *Mtb* growth and positively affect the outcome of the infection, by limiting steady-state bacterial infection in the lungs (the bacterial 'set-point'). In no case, however, has immunization prevented infection or achieved sterilizing immunity. These data suggest that pre-exposure vaccination strategies will be successful in limiting early *Mtb* colonization, but they may not succeed in preventing infection.

It is also apparent that prior *Mtb* infection does not generate highly effective immunity to reinfection. A retrospective examination of a number of human studies has shown that *Mtb* reinfections are not uncommon, suggesting that at least some humans do not generate protective immunity following infection (163, 164). In mice, this question has been addressed experimentally following antibiotic treatment of infected animals (28, 165). *Mtb* infection of drug-treated mice resulted in earlier IFN production by effector T cells, and the treated mice exhibited a lower the bacterial set-point (126, 152). Another recent study has shown that antigen-specific memory phenotype CD8<sup>+</sup> T cells were present in lungs, LNs, and spleen 12 weeks after antibiotic treatment, suggesting that memory T cells are generated during *Mtb* infection (166). Nevertheless, the apparently relatively poor protective value of prior *Mtb* infection has led some investigators to question whether effective vaccination is attainable against TB (152, 164).

Why is it, then, that neither immunization nor prior infection has been able to achieve a better outcome? One possibility, discussed above, is that *Mtb*, once established in host cells, is impervious to defenses triggered by T cells. This would explain why early T-cell responses can sometimes limit but cannot prevent infection. Another possibility is that natural infection and/or vaccination does not generate effective T-cell memory. *Mtb* infection may induce abortive memory T-cell response, perhaps due to inflammatory conditions that do not favor memory generation. Alternatively, vaccination strategies attempted thus far may not have succeeding in generating sufficient numbers of memory T cells sufficient to elaborate effector functions that will control TB. In this regard, correlates of protective T-cell-mediated immunity are unavailable in either mice or humans that would allow us to evaluate what constitutes a protective memory response. Thus, much more work will be required to first understand and then to harness early memory T-cell responses during both primary and secondary infections.

#### Conclusions

Available evidence indicates that early innate and adaptive responses to *Mtb* are of critical importance for successful containment of infection. These events are poorly understood, however, in part because low-dose aerogenic inoculation does not trigger a robust early inflammatory response and because very early T-cell responses have been difficult to study. Important questions remain regarding whether *Mtb* delays the onset of early immunity, either by accident or design, and whether true memory T lymphocytes can be elicited by vaccination. New tools are now available to address these questions and have the potential to improve our understanding of TB immunity.

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# **Fig. 1. Schematic of early T-cell responses after aerosol** *Mtb* **infection** Idealized representations of T-cell priming, effector cell generation, and proliferation, in three different tissues, are shown.



#### Fig. 2. Detection of transgenic CD4<sup>+</sup> T cells in the lungs

Transgenic IA<sup>b</sup>/ESAT-6-specific CD4<sup>+</sup> T-donor cells were visualized in a mouse lung on day 17 post-aerosol inoculation. The transgenic CD4<sup>+</sup> T cells (yellow arrows) were detected using an anti-CD90.2 antibody (visible in red). The cell nuclei were stained using DAPI. The image was captured using a Zeiss fluorescence microscope at  $\times$  40 magnification.