

Host Innate Immune Response to *Mycobacterium tuberculosis*

KAMLESH BHATT¹ and PADMINI SALGAME^{1,2}

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This review focuses on recent progress in our understanding of *Mycobacterium tuberculosis* survival in macrophages, the interaction of *M. tuberculosis* with Toll-like receptors (TLRs) and the establishment of the link between innate and adaptive immunity, and TLRs and interferon- γ -mediated antimicrobial pathways in macrophages. We also propose a paradigm that TLR2 signaling regulates the magnitude of the host Th1 response leading to either *M. tuberculosis* persistence and latent infection or replication and disease.

KEY WORDS: *Mycobacterium tuberculosis*; innate immunity; TLR2; dendritic cells; macrophages.

INTRODUCTION

The naissance of innate immunity was the description of macrophage phagocytosis by Metchnikoff (1). Today, the science of innate immunity is more than just phagocytosis. The innate immune response comprises several different cell types, has its own receptor system to recognize the presence of pathogens, and is a key to the initiation of an adaptive immune response in the host. No wonder, successful pathogens have evolved ways to evade innate immune killing in order to find a niche in the host. In this review, we will discuss host innate immunity generated in response to the pathogen, *Mycobacterium tuberculosis* (Mtb).

Mtb infects approximately one-third of the world's population (2). Close to eight million new cases of tuberculosis occur each year, accounting for approximately 7% of all deaths and 26% of all avoidable adult deaths in developing countries (3). Despite the implementation of TB control programs, case rates continue to soar where

the prevalence of HIV infection is high (4). The situation is further complicated by a worldwide increase in drug resistant and MDR-TB, and the recent reports of XDR TB (5). Thus, the resurgence of TB truly constitutes a global health crisis (6).

Tuberculosis begins with the inhalation of Mtb-containing aerosols into the pulmonary alveoli. Here, the bacteria bind to phagocytic receptors and enter resident alveolar macrophages, dendritic cells, and monocytes recruited from the bloodstream. Besides expressing phagocytic receptors, macrophages and dendritic cells also express Toll-like receptors (TLRs) that recognize conserved molecular patterns expressed on pathogens (7–9). Ligation of TLRs by these pathogen-specific ligands initiates a signal transduction pathway in the host cell that culminates in the activation of NF κ b and the induction of cytokines and chemokines (10) that are crucial to eliciting the adaptive immune response against the pathogen. Consequently, activation of TLR is an important link between innate cellular response and the subsequent activation of adaptive immune defense against microbial pathogens.

As depicted in Fig. 1, a small percentage of individuals, despite exposure to Mtb, remain uninfected, most likely due to the expression of high innate immunity. However, in the majority of individuals who are exposed to Mtb, the innate response cannot protect from infection, and effector Th1 cytokines of the adaptive immune response are necessary to restrict bacterial growth and mediate protection. The adaptive immunity generated in these people, although protective, nonetheless does not induce sterilizing immunity. These individuals therefore remain latently infected, and are vulnerable to disease reactivation when their immune surveillance weakens or when their immune response is compromised. Reactivation tuberculosis contributes significantly to the morbidity and mortality associated with the disease (11, 12), and is believed to account for a substantial portion of TB cases in HIV-infected individuals (13). In another small proportion of individuals, infection leads directly to primary tuberculosis due to a

¹Department of Medicine, Centre for Emerging Pathogens, UMDNJ-New Jersey Medical School, MSB A902, 185 South Orange Avenue, Newark, New Jersey, 07101.

²To whom correspondence should be addressed to; e-mail: salgampa@umdnj.edu.

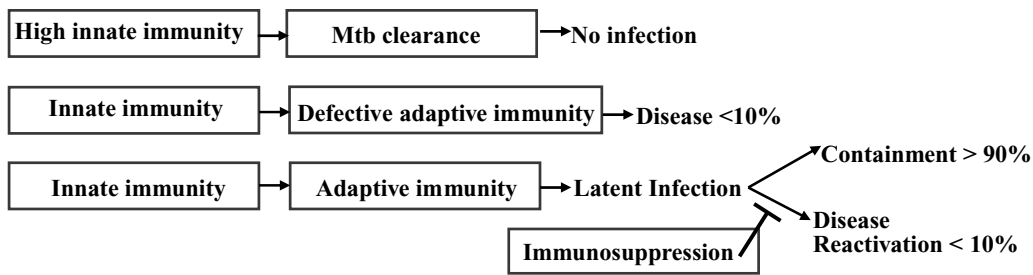


Fig. 1. Three possible outcomes of Mtb exposure. In some individuals, Mtb is eliminated by the host immediately upon inhalation. The frequency and the cause of such spontaneous healing are not certain. In the second and the largest group of individuals, infection is contained as a result of successful granuloma formation, a function of strong innate and adaptive immune response by the host, which results in latent infection. In this group, reactivation of latent infection can occur due to factors such as aging or the immunocompromised status of the host. In a small number of infected individuals, adaptive immunity fails and they develop primary tuberculosis.

failure of their adaptive immune response to control the initial bacterial replication.

In the last several years, we have gained a better appreciation regarding host innate immune response to Mtb, although much needs to be learned regarding the nature of the innate response that prevents establishment of Mtb infection. This review focuses on our current understanding of the regulation of host innate immune response to Mtb and how it interfaces with the adaptive immune response. The review is not all-inclusive, and only areas where substantial progress has been made will be addressed. Specifically, we will discuss (i) Mtb entry and subsequent survival inside macrophages; (ii) Mtb interaction with innate receptors, specifically TLRs; (iii) Mtb-induced cytokine and chemokine induction in macrophages and dendritic cells; (iv) maturation and migration of dendritic cells—a key step that links innate and adaptive anti-Mtb immunity; (v) modulation of innate immunity by effectors of the adaptive immune response. Finally, in closing, we will explore the evolving paradigm that innate immune response generated in the host in response to Mtb infection is not only important for initiating anti-Mtb immunity, but the innate response concomitantly also activates regulatory pathways in the host. Whether the regulatory mechanisms activated are important for controlling the magnitude of the host immune response to prevent immunopathology, or whether this is a virulence strategy for the TB bacillus to persist in the host will be discussed.

MTB PHAGOCYTOSIS AND SURVIVAL INSIDE MACROPHAGE

Macrophages lining the alveolar spaces of the lungs represent the first line of defense upon aerosol infection of the host with Mtb. *In vitro* studies have implicated comple-

ment receptor (CR)3 as a major receptor on macrophages for phagocytosis of Mtb (14). Nonetheless, several other macrophage surface receptors, such as CR1, CR4, Mannose receptor, CD14, and Scavenger receptors can also recognize and bind Mtb *in vitro* (15). Pulmonary surfactant protein A (Sp-A) too enhances Mtb uptake by human macrophages (16, 17). In this context, it is worth noting that although CR3 was determined as the primary mode of macrophage entry for Mtb *in vitro*, mice lacking CR3 exhibited similar bacterial burden and host response to that of CR3-sufficient mice (18). Cholesterol accumulates at the site of Mtb entry into macrophages, and depleting cells of cholesterol prevent Mtb internalization (19). This indicates that perhaps cholesterol accumulation around phagocytic receptors rather than the nature of the receptor itself dictates Mtb uptake. Whether *in vivo* a similar relocation of cholesterol occurs at sites of Mtb entry into macrophages and whether it provides a distinct advantage to the TB bacillus remains to be determined.

Once inside the host cell, Mtb successfully evades destruction by the innate microbicidal machinery. Armstrong and Hart in papers published in the early 1970s (20, 21) shaped our understanding of how Mtb might persist in the host. Their work showed that Mtb vacuoles did not fuse with the lysosomal compartment. Substantive work from several laboratories has built on this seminal observation to provide a detailed insight into the molecular events that arrest the maturation of the Mtb phagosome and prevent its further biogenesis and acquisition of lysosomal components. The early trafficking pattern of the Mtb phagosome is normal and exhibits fusion with certain early endosomal compartments, since both iron (22, 23) and glycosphingolipids (24) are found associated with the Mtb phagosome. The arrested phagosome, however, lacks the vacuolar ATPase and lysosomal hydrolases (25). Subsequent studies that monitored the trafficking pattern

of the Mtb phagosome showed that the arrest occurs between the acquisition of the endocytic vesicles Rab5 and Rab7. The Mtb phagosome is associated with Rab5 (26), but not Rab7 (27), and furthermore exhibits reduced recruitment of the early endosomal autoantigen 1 (EEA1) (28), an effector molecule of Rab5 required for organelle tethering and delivery of lysosomal hydrolases, cathepsins, and vacuolar ATPases from the trans golgi network to the phagosome. The Mtb phagosome also lacks a specific type III phosphatidylinositol 3-kinase, hVPs34 (29) whose activation product phosphatidylinositol 3-phosphate aids the retention of EEA1 to the endosomal membrane (29).

How and what components of Mtb block the Mtb phagosome from undergoing the typical phagosome biogenesis? The arrest of the Mtb phagosome at the Rab 5 stage and its inability to proceed through the maturation pathway, at least partly, results from Mtb-induced inhibition of sphingosine kinase activity and subsequent Ca^{2+} signaling pathway in the cell, a step necessary for recruitment of hVPs35 to membranes of organelles (30). Another study reported that maturation of the Mtb phagosome is impeded because Mtb suppresses phagosomal actin assembly (31). Yet another study determined that the Mtb phagosome arrest was dependent on its initial fusion with early endosomes and acquisition of iron (32). Manosylated lipoarabinamannan (ManLAM), the Mtb analog of host phosphatidylinositol-3 phosphate is responsible for actively inhibiting Mtb phagosome from fully maturing and acquiring lysosomal hydrolases (33, 34). Indeed, it has been shown that phagocytosis of Mtb via binding of its cell surface LAM to the mannose receptor on human macrophages led to the non-fusogenic phenotype of the Mtb phagosome (35). Interestingly, other Mtb lipids such as PIMs enhance Mtb phagosome fusion with early endosomes, possibly providing the phagosome access to host nutrients (36). Besides lipids, a protein from Mtb, protein kinase G (PknG) has also been implicated in interfering with the transfer of Mtb phagosome to the lysosomal compartment (37).

Our understanding of Mtb phagocytosis, the biogenesis of Mtb phagosome, and intracellular growth of Mtb is almost entirely derived from studies examining interaction of Mtb with macrophages. However, it is necessary to also understand the handling of Mtb inside dendritic cells, since in response to an aerosol challenge with Mtb, dendritic cells take up Mtb, and as will be discussed later, are crucial to linking the innate and adaptive immune responses. Besides the expression of CR3 and mannose receptor, dendritic cells are endowed with an additional phagocytic receptor for binding Mtb. Dendritic cells express the C-type lectin, DC-SIGN (DC-specific intercel-

lular adhesion molecule-grabbing nonintegrin), and Mtb can bind DC-SIGN through manLAM expressed on their surface. A comparative analysis of Mtb survival within human macrophages and dendritic cells revealed that unlike macrophages, dendritic cells did not support intracellular growth of Mtb (38, 39). Despite being a key player in the innate response to Mtb, we know very little regarding the trafficking of Mtb vacuole inside dendritic cells; except for a study that reported that endosomal trafficking is significantly reduced compared to that in macrophages (39). Clearly, more detailed studies examining the intracellular fate of Mtb inside dendritic cells are needed.

It must be emphasized that Mtb replication in the macrophage is also controlled at the level of host factors. Expression of a candidate gene *IntracellularPathogenResistance* (*lpr1*) within the *sst1* locus limits Mtb multiplication in the host (40). Variants in the human equivalent of the *lpr1* gene SP110 were shown to be associated with genetic susceptibility to TB in a study of families in West Africa (41). Another association study in human TB, also in West Africa, however, found no association of human pulmonary TB with SP110 variants (42). Undoubtedly, more studies in genetically different populations are needed to equivocally determine the role of SP100 and other potential candidate genes in susceptibility to TB.

MTB INTERACTION WITH TLRs

Engagement of TLR by Mtb ligands is an early event in the interaction of Mtb with its host cell. Accumulating data indicate that Mtb expresses a large repertoire of TLR2 ligands. The 19-kDa lipoprotein (LpqH), a secreted antigen of Mtb, was the first Mtb ligand shown to interact specifically with TLR2 to induce TNF α and nitric oxide production from both murine and human macrophages (43). In addition, the 19-kDa lipoprotein is a major inducer of interleukin (IL)-12 production in human monocytes (43). LprA (Rv1270) (44) and LprG (Rv1411c) (45) are two other mycobacterial lipoproteins that are TLR2 agonists. In addition to lipoproteins, lipomannan (46) and phosphatidyl-*myo*-inositol mannoside (PIM) (47, 48) also interact with TLR2 to initiate cellular activation (48). However, with regards to PIMs, Abel *et al.* (49) demonstrated that PIM structures can also elicit cellular activation via TLR4. They showed that PIM was able to induce NF κ B activation in a dose-dependent manner in stable TLR4 and MD-2 Ba-F3 transformants. A systematic biochemical characterization of four acyl forms of lipomannan (LM) from *M. bovis* BCG that differed in their degree of acylation, indicated that only the triacylated LM was a potent TLR2 agonist (50).

Interestingly, ManLAM derived from virulent Mtb fails to activate either TLR2 or TLR4-transfected cells (51). In contrast, AraLAM purified from fast-growing mycobacteria is capable of TLR2-mediated cellular activation (51).

Studies aimed at determining the requirement of TLR4 in controlling Mtb infection following an aerosol challenge showed that lack of TLR4 did not compromise host resistance to TB (52, 53). However, a high dose of Mtb infection did lead to enhanced susceptibility in the absence of TLR4 signaling (49). It is interesting that despite a large collection of TLR2 agonists on the TB bacillus, murine studies indicate that TLR2 is not essential for host resistance against tuberculosis. In a model of low-dose aerosol infection, TLR2 (53, 54) deficiency did not affect host defense against Mtb infection. However, in one of the two studies (53) with high-dose aerosol infection, a role for TLR2 in host resistance was revealed. The TLR2-deficient mice were not compromised in their ability to induce Th1 immunity, but on the contrary, exhibited exaggerated immunopathology. *In vitro* studies have shown that engagement of TLR2 with Mtb ligands induces inhibition of macrophage MHC class II antigen presentation (55) and also blocks macrophage responsiveness to IFN γ (56, 57). Together with the *in vivo* studies, these *in vitro* findings that TLR2 signaling negatively modulates macrophage functions point to the need for future studies designed to examine whether the negative signaling from TLR2 curtails Th1 activation, and whether this is important for balancing protection and immunopathology in the host.

Our studies examining the *in vitro* interaction of TLR with live Mtb reported that in response to Mtb, dendritic cells secreted copious amount of IL-12, while the secretion was limited in infected macrophages. The study also reported that Mtb induced rapid and significantly higher remodeling at the IL-12p40 promoter in dendritic cells in comparison to macrophages. The mechanism behind the differential remodeling at the IL-12p40 promoter and subsequent IL-12 release was shown to be due to differences in TLR usage by macrophages and dendritic cells. Mtb induced IL-12 secretion from dendritic cells in a TLR9-dependent manner while in macrophages it was TLR2-dependent (58). Consistent with this, the greatest effect on the progression of tuberculosis disease was seen in mice doubly deficient in TLR2 and TLR9 (59).

Although IFN γ is undoubtedly necessary for resistance against Mtb infection (60), it is of interest that there exist antimycobacterial pathways that are independent of IFN γ and are induced by TLR in human macrophages. For example, it has been known for some time that activation

of the Vitamin D3 pathway controls Mtb replication in human macrophages (61). Also, it had been documented that Vitamin D deficiency is a risk factor for tuberculosis (62). Only recently, however, Modlin and colleagues deciphered the mechanism for Vitamin-D3-mediated antimicrobial pathway. They demonstrated that TLR2-mediated activation of macrophages upregulated the expression of Vitamin D receptor and Vitamin-D-1-hydroxylase genes, leading to the induction of the antimicrobial peptide, cathelicidin (63). The study from Modlin's group also showed that African American individuals who are more susceptible to Mtb infection and disease were not efficient in inducing the antimicrobial peptide, cathelicidin. The TLR2-mediated innate mechanism of mycobacterial killing provides a scientific basis for tuberculosis treatment of a century ago: exposure to sunlight. Other TLR-induced killing mechanisms may also participate in the innate response. For instance, CpG, an activator of the TLR9-pathway, also induces rapid antimycobacterial responses in macrophages, in a phospholipase D-dependent manner (64). These innate mechanisms for killing Mtb provoke future investigations of whether individuals who never become infected with Mtb have the capacity to activate these pathways and overpower the Mtb-induced block in phagosome maturation.

MTB-INDUCED UPSURGE OF CYTOKINE AND CHEMOKINE SECRETION

A major consequence of Mtb interaction with the TLRs on macrophages and dendritic cells is the burst in cytokine and chemokine secretion. The induction of these effector molecules regulates the formation of the granuloma and is responsible for initiating and shaping the adaptive immune response to Mtb. The contribution of adaptive immunity to the evolving tubercle granuloma in the lung will not be elaborated in this review. The reader is referred to other reviews on the topic (65–67).

Cytokines Important for Induction of Th1 Immunity

Clearly, induction of cellular Th1 immunity is critical for protection against tuberculosis as evidenced by enhanced disease in the HIV-infected (68) and from experiments of nature where individuals carrying defective genes for IFN γ R and IL-12R (69) are exquisitely susceptible to intracellular pathogens, including mycobacteria. Currently, there are three well-defined cytokines that steer naïve T cells toward Th1 commitment (70). IL-12 was the first cytokine to be described with potent Th1 promoting attributes, followed by the discovery of IL-23

(shares the p40 component with IL-12) and the recent addition of IL-27 to this list. Work from several laboratories has revealed that the three cytokines together orchestrate Th1 responses, with IL-12 being the prototypic and dominant cytokine that affects both the induction and maintenance of Th1 immunity. IL-23, on the other hand, has activities on memory T cells and IL-27, secreted prior to IL-12 by antigen-presenting cells, is involved in Th1 initiation.

In patients with tuberculosis pleuritis, a clinical form of disease that is mostly self-healing, high IL-12 levels were found in the pleural fluid (71). Two studies comparing murine macrophages and dendritic cells demonstrated that dendritic cells release significantly higher IL-12 than did macrophages in response to live Mtb (72, 73). *In vitro*, Mtb-infected dendritic cells also primed naïve T cells toward Th1 development, while macrophages did not; though, IL-23-secreting macrophages were capable of inducing Mtb-specific Th1 response (74).

Early studies in the murine model of tuberculosis clearly demonstrated that the cytokine IL-12 that is necessary to drive Th1 responses and IFN γ —the effector molecule of the Th1 response—were both necessary for protection against Mtb infection. Mice deficient in the p40 component of IL-12 or in IFN γ (GKO) were both highly susceptible to Mtb infection. Exogenous supplementation of IL-12 at the onset of disease led to reduction in bacterial burden and delayed the lung pathology in the relatively susceptible Balb/C strain of mice (75). However, IL-12 supplementation did not lead to enhanced protection in GKO mice indicating that IFN γ is downstream of IL-12 and is the effector molecule mediating protection in the host. In another study (76), the role of endogenous IL-12 was studied by neutralization with anti-IL-12 antibodies. It was found that in Balb/C mice neutralization of IL-12 at the onset of infection led to disruption in the ability of the host to contain infection; however, neutralization of the cytokine after the onset of infection (third week) did not affect bacterial replication. This suggests that the presence of IL-12 is more critical during early infection when anti-Mtb adaptive immunity is being shaped toward Th1-type. However, in later studies, reconstitution of IL-12-p40 gene-deficient mice with recombinant IL-12 only during the early phase of infection was determined not to be sufficient to provide long-term immunity, despite early control of bacterial growth. Transfer of immune CD4 T cells from Mtb-infected wild-type mice to Rag $^{-/-}$ provided immunity against infection. However, similar reconstitution of immune CD4 T cells into Rag $^{-/-}$ mice that were also deficient in IL-12-p40 failed to induce protection. This provides experimental evidence that sustained production of IL-12 throughout the course of

infection is necessary to maintain antibacterial immunity in the host (77). A reason why this study differed from the previous where IL-12 was shown not to be necessary for long-term immunity may probably be due to the presence of residual IL-12 activity in the neutralization experiments.

To further characterize whether susceptibility to Mtb infection in the absence of p40 is due to the lack of biologically active IL-12 or is a consequence of defective IL-23 production, mice deficient in the specific components of IL-12 and IL-23, p35 and p19, respectively, were studied. Mice lacking p19 were able to control Mtb infection as well as the wild-type mice (78, 79). Mice lacking p35 were able to control bacterial replication slightly better than the p40-deficient mice (79, 80), but in comparison to the p19 knockout mice exhibited significantly higher bacterial burden. Mice doubly deficient in p35 and p19 genes were as susceptible as the p40-gene-deficient mice (79). Exogenous delivery into the lung of IL-23 via adenoviral vectors enhanced anti-Mtb immunity, upregulated IL-17 expression, and reduced bacterial burden in the lungs of Mtb-infected mice (81). Together, these data indicate that IL-23 is less critical for protection against Mtb, and only provides a moderate level of protection to the host in the absence of biologically active IL-12. On the other hand, IL-12 has a far more vital role in the generation of protective anti-Mtb immunity.

IL-27 is also an IL-12-related cytokine and WSX-1 is a component of the IL-27R complex (82). IL-27/IL-27R signaling, interestingly, exhibits both pro- and anti-inflammatory properties. Infection of WSX-1 $^{-/-}$ mice with Mtb revealed that, in the absence of IL-27R signaling, there was a reduced bacterial burden accompanied by enhanced CD4 infiltration into the lungs (83). Another group examining Mtb infection in the same WSX-1 $^{-/-}$ mice observed increased IL-12-p40 and TNF α expression and enhanced IFN γ production from CD4 T cells. This group also reported reduced Mtb burden in the lungs of infected WSX-1 knockout mice in comparison with wild-type mice. Despite restricted bacterial growth, the WSX-1 knockout mice succumbed to infection due to exaggerated immunopathology, a scenario similar to what was first reported with *Toxoplasma gondii* infection in this strain of knockout mice. Under conditions where pathogens do not induce a Th1 response in the host, for example in *Leishmania major* infection of Balb/c mice, absence of WSX-1 resulted in the generation of protective Th1 response with concomitant Th2 downregulation in the host (84). A tenet for future perusal is that the anti-inflammatory activity of IL-27 is perhaps more critical than its Th1 promoting activity in response to pathogens that have high Th1-inducing potential (85, 86).

TNF

TNF α plays an important role in regulating the pathology of tuberculosis (87). TNF α exists in both soluble and membrane bound forms and signals through TNF α R. Mtb infection leads to TNF α secretion by macrophages, dendritic cells, and T cells (60). Secretion of TNF α by Mtb-infected macrophages is a potent mechanism to induce killing of Mtb via generation of reactive nitrogen intermediates in conjunction with IFN γ (88). An attribute of membrane TNF α is to induce apoptosis of the Mtb-infected alveolar macrophages (89), and thereby indirectly contribute to the reduction of bacterial burden. TNF α 's ability to induce alveolar macrophage apoptosis may also be important in the cross-presentation of Mtb antigens for CD8 cytotoxic T cell priming (90). It has also been suggested that inhibition of TNF α -mediated macrophage apoptosis is a virulence strategy of Mtb. Avirulent H37Ra induced TNF α -dependent macrophage apoptosis, while virulent H37Rv released soluble TNFR2 that reduced TNF α activity and subsequent apoptosis of macrophages (91). Although it needs to be examined in more detail, TNF α has been shown to support the growth of Mtb in human monocytes and macrophages (92).

The requirement for TNF α in host defense against Mtb infection was demonstrated in studies which showed that mice treated with antibody to TNF α became more susceptible to BCG infection and exhibited malformed granulomas (93). Mtb infection of mice lacking TNF receptor or neutralization of TNF α activity in mice also led to the failure to control bacterial replication resulting in enhanced susceptibility (94). This study indicated that TNF α contributed to maintaining host resistance by inducing the production of reactive nitrogen intermediates by macrophages. Later studies have indicated that TNF α also participates in setting the chemokine circuitry in the developing granuloma. Mice lacking TNF α had reduced chemokine expression in lung granulomas (95–97) and this resulted in reduced T cell infiltration into the lungs and a failure to form a productive granuloma.

For the most part, the immunological forces that control reactivation remain ill defined, except for the knowledge that TNF α is a major player (98). Studies in a murine latent model of tuberculosis from the Chan and Flynn laboratories clearly demonstrated that neutralization of TNF α during the latent/persistent phase induced reactivation in C57BL/6 mice, as indicated by the enhanced bacterial burden in the lungs (99). Further, histological examination of lung tissue from TNF α -neutralized animals revealed a disorganized granuloma with indications of a lack of cellular turnover and increased fluid accumulation. NOS2 expression was attenuated, while IL-10 expression was

upregulated in the lungs. Immunohistochemical analysis indicated an increased presence of apoptotic T cells and macrophages in the lung, a feature not seen previously in other reactivation models. The importance of TNF in maintaining Mtb in a chronic/persistent phase in mice has been corroborated in humans. It has been observed that the use of anti-TNF α antibody in patients undergoing treatment for rheumatoid arthritis has resulted in reactivation of tuberculosis in some latently infected individuals (100, 101).

Mtb-Induced Chemokines

In vitro and *in vivo* studies provide evidence for the participation of chemokines in the control of TB. It is present in the innate and adaptive immune response to Mtb (96, 102). Mtb infection of both human and murine macrophages results in the secretion of a large number of chemokines, including CCL2, CCL3, CCL7, CCL12, CXCL2, and CXCL10 (96). A comparative chemokine expression analysis showed that lung interstitial macrophages from a susceptible mouse strain expressed significantly high levels of CXCL13 and CXCL14, while higher expression of CXCL9 and CXCL0 was found in macrophages from the resistant strain of mice (103). Regulation of chemokine production in macrophages is predominantly regulated by TNF α . Mtb infection of macrophages leads to the production of TNF α which, in turn, regulates the secretion of a plethora of chemokines from macrophages, including CCL2, CCL3, CCL4, CCL5, CXCL10, and CXCL13 (104). Mtb-infected dendritic cells also secrete chemokines, including CXCL9, CXCL10, CCL3, and CCL4. CXCL10 secretion was IFN α -dependent and in conjunction with CXCL9 and CXCL3 acted to recruit inflammatory cells to the site of Mtb infection (105).

Studies to examine the role of chemokine and chemokine receptors in host resistance against Mtb infection has led to conflicting results, in great part due to redundancy in the function of chemokines and their receptors. In a mouse model of tuberculosis, it has been shown that the first step in recruitment of cells into the lung, specifically recruitment of immature dendritic cells and monocytes to the site of infection, is mediated by CCR2 and, as a consequence, CCR2 $^{-/-}$ mice (106) are more susceptible to Mtb infection. These mice also show defective recruitment of dendritic cell to the draining lymph nodes resulting in delayed and reduced priming of naive T cells. However, a later study (107) indicated that susceptibility of CCR2 knockout mice to Mtb infection was dose dependent. As seen with high dose infection, a low-dose aerosol challenge of the CCR2 knockout mice with

Mtb also resulted in reduced cellular migration to the lungs and delayed priming. However, there was no change in bacterial burden or susceptibility to infection. Using chimeric mice, where either the myeloid or the lymphoid compartment was lacking CCR2, it was determined that expression of CCR2 on macrophages and dendritic cells was important for the recruitment of T cells to lungs (108). However, mice deficient in CCL2/MCP-1, which is a ligand for CCR2, do not show reduced susceptibility to Mtb infection, thereby indicating that *in vivo* other chemokines such as CCL7, CCL8, and CCL12 can compensate for the lack of CCL2/MCP-1 (109).

In addition to its expression on granulocytes and macrophages, CCR5 is also present on immature dendritic cells and Mtb modulates its expression. Indeed, mycobacterial Hsp70 can interact with CCR5 on immature dendritic cells and induce their maturation and the interaction also induces IL-12 secretion from dendritic cells (110). Despite CCR5-mediated IL-12 production and the enhanced production of CCR5 ligands, MIP-1 α , MIP-1 β , and RANTES in the lungs of Mtb-infected mice, absence of CCR5 did not affect the ability of the host to control bacterial replication in the lung (111, 112). The latter study, in addition, observed that CCR5 $-/-$ mice exhibited increased bacterial burden in the draining lymph nodes (112). The intriguing possibility that CCR5 signals impede Mtb-bearing dendritic cell migration resulting in enhanced accumulation of Mtb in the draining lymph nodes needs further scrutiny.

CCR7 expression on cells guides their migration to the draining lymph nodes where its cognate ligands CCL19 and CCL21 are present. Indeed, Mtb infection upregulates CCR7 expression on dendritic cells, but absence of CCR7 did not enhance bacterial replication (113). Although it must be noted that the granulomas of CCR7 $-/-$ mice had altered granuloma architecture with enhanced inflammation and a lack of follicular B cell architecture. How these changes in the granuloma affect host resistance is still not clear. As discussed in the next section, Mtb-infected dendritic cells migrate to the draining lymph nodes to initiate an immune response. Therefore, it would be important to determine what chemokine/receptor gradient controls the migration of Mtb-infected dendritic cells from lungs to draining lymph nodes.

CXCR3 is a chemokine receptor preferentially expressed on activated Th1 cells and regulates their migration in response to ligands CXCL10, CXCL9, and CXCL11. Given the importance of Th1 in host resistance against TB, C57BL/6 mice deficient in CXCR3 were studied following Mtb infection. Despite the ability of CXCR3 to regulate the migration of Th1 cells, absence of the receptor did not affect Mtb replication in the host. The

CXCR3 knockout mice, however, did exhibit a neutrophil deficit in the granuloma. The consequence to host resistance of reduced neutrophils in the granuloma remains unclear (114). Contrary findings were reported in a recent study that examined mice lacking CXCR3 on the BALB/c background. In this study, the CXCR3-deficient mice exhibited heightened resistance to Mtb infection in the chronic phase when compared with wild-type mice (115), and the mice also had enhanced T cell activation (115). The authors of the paper suggest that enhanced resistance in BALB/cCXCR3 knockout mice could have resulted from the absence of the immunosuppressive CXCR3–CXCL10 chemokine gradient. Certain chemokine gradients, including CXCR3–CXCL10, have recently been recognized as immunosuppressive and to interfere with the formation of the immunological synapse (116). Together, these studies highlight the emerging recognition that the chemokine circuitry activated during Mtb infection may not only regulate cellular recruitment, but also directly impact on the function of immune cells. The studies also highlight the role of genetic differences in regulating chemokine functions.

IL-10

Dendritic cells and macrophages in response to Mtb produce the immunosuppressive and anti-inflammatory cytokine IL-10. Interestingly, dendritic cells secrete substantial IL-12 in response to Mtb infection and can prime naive T cells toward Th1-type, despite concomitant secretion of IL-10 (73, 117). IL-10-secreting CD8 suppressor/regulatory T cells are associated with susceptibility to Mtb infection (118) and T cells expressing both IFN γ and IL-10 have been isolated from the bronchoalveolar lavage fluid of TB patients (119). Additionally, depressed T-cell IFN γ responses in pulmonary tuberculosis was shown to be associated with the induction of IL-10 from monocytes (120). Interestingly, in patients with pleural TB, considered as the resistant and self-healing form of the disease, IL-10 is found along with IFN γ at sites of infection in the pleural fluid (121). *In vitro*, IL-10 downregulates the production of IL-12 in human monocytes infected with Mtb (122). Also, IL-10 down modulates the activity of CD4 and CD8 T cells via downregulation of costimulatory molecules on macrophages (123). In addition, IL-10 inhibits the proliferation of IFN γ producing T cells and $\gamma\delta$ T cells (124). Although absence of IL-10 did not enhance resistance to Mtb infection in IL-10 knockout mice (125), transgenic over-expression of IL-10 resulted in reactivation of chronic disease (126). Similarly, expression of human IL 10 in mice under the control of MHC II promoter enhanced the susceptibility to disease, independent of

T-cell-derived IL-10. The Mtb-infected macrophages from these transgenic mice exhibited reduced antimycobacterial capacity (127). That IL-10 may have a role in TB is suggested by the fact that polymorphism in murine SLC11 A1, a tuberculosis susceptibility locus, has been associated with variation in IL-10 production (128). Together, these data suggest that IL-10 is induced by Mtb and suppresses the generation anti-Mtb immunity. However, Th1 cytokines are often found along with IL-10. Perhaps the relative quantities of the two cytokines determine whether Th1 immunity is suppressed or not.

DENDRITIC CELL MATURATION, MIGRATION, AND ANTIGEN PRESENTATION

Recent work from several laboratories has focused on dissecting the role of dendritic cells in Mtb infection. Upon its interaction with Mtb, dendritic cells undergo a repertoire of phenotypical changes, a process termed as maturation. This process, which is TLR-dependent, brings forth three major phenotypic changes in dendritic cells: upregulation of costimulatory molecules—CD40, B7.1 and B7.2, heightened expression of adhesion molecules, and upregulation of chemokine receptor—CCR7. Whereas immature dendritic cells are efficient at Mtb phagocytosis and exhibit enhanced microbicidal property, maturation endows them with the role of an efficient antigen presenter and initiator of adaptive immune responses (129).

Following Mtb phagocytosis and concomitant TLR activation, the next step in the development of host immunity is the transport of pathogen from the lung to the draining lymph nodes, where the matured dendritic cells can present antigen to naive T cells and initiate the process of adaptive immune response. Although Mtb uptake and engagement of TLR signaling for cellular activation occurs in macrophages and dendritic cells, only the latter cell type was shown to acquire the capacity to upregulate CCR7 expression and migrate to draining lymph nodes (130). Consistent with this study that tracked intratracheally instilled cells, endogenous lung dendritic cells also exhibited similar migratory property. Following intratracheal infection of mice with GFP-expressing BCG, dissemination of mycobacteria from the lung was initiated by the migration of infected dendritic cells to the draining lymph nodes (131), despite predominant infection of alveolar macrophages. Another study in BCG-infected mice also demonstrated that dendritic cells, and not macrophages, were the antigen-presenting cells responsible for priming naive T cells (132). Direct evidence for the role of dendritic cells as the priming APC for initiating pulmonary immunity came from the study where

mice depleted of CD11c + dendritic cells exhibited delayed CD4 responses to Mtb and worsening disease (133). It has been argued that in addition to ferrying antigen for T cell priming, migration of dendritic cells to the lymph nodes may also aid in Mtb dissemination (131).

Are other cell types involved in transporting Mtb antigens to the draining lymph nodes? Indeed, a recent report implicates neutrophils as the carrier of live BCG following intradermal vaccination from peripheral tissue to the DLN capsule (134). Whether neutrophils participate in antigen transport during a pulmonary infection with Mtb would be worth investigating, particularly since neutrophils appear to marginally influence early immune responses and the architecture of the ensuing granuloma (135, 136). Following Mtb infection, there is an influx of macrophages and dendritic cells from the periphery into the lung. The relative contribution of interstitial versus the newly recruited dendritic cells in Mtb transport from lung to the draining lymph nodes is also not clear. However, absence of CCR2 was shown to impair macrophage and dendritic cells trafficking into the infected lungs resulting in susceptibility to Mtb infection (108), suggesting that dendritic cells recruited to the lung may also function to carry Mtb to the draining lymph nodes.

Collective data indicate that dendritic cells are the antigen-presenting cells that migrate to the draining lymph nodes, and process and present Mtb antigens on MHC Classes I and II to naive CD4⁺ and CD8⁺ cells, respectively. This review will not address the mechanisms of antigen processing and presentation and other molecular events controlling T cell activation. However, we discuss here one study from Kaufman's group that demonstrated a detour pathway for how antigens of Mtb, presumably confined within the phagosome, are delivered to Class I molecules. This study demonstrated that dendritic cells take up apoptotic vesicles containing Mtb, the vesicles are then degraded in an endosomal-dependent manner and Mtb antigens are cross-presented on MHC Class I molecules to CD8 T cells (137).

Since maturation and migration of dendritic cells is such a key step in linking innate and adaptive immunity, it is not surprising that Mtb negatively interferes with this step. IL-1 β release from mycobacteria-infected antigen-presenting cells inhibits dendritic cell maturation (138) and virulent Mtb has been reported to impair the maturation of monocyte derived dendritic cells (139). The migration of dendritic cells appears to be regulated to some degree by IL-12p40 homodimers (140). Whether IL-10, which can impede dendritic cell migration (141), functions by downregulating the p40 homodimers is worth considering. In addition, one also needs

to examine whether in human infection *Mtb* interference with the dendritic cell maturation and migration process is dependent on the degree of virulence of the infecting clinical strain and its ability to induce IL-10 production.

MODULATION OF INNATE IMMUNITY BY EFFECTORS OF THE ADAPTIVE IMMUNE RESPONSE

In effect, despite the early induction of chemokines and cytokines from macrophages, *Mtb* is able to skillfully avoid the innate antimicrobial defense mechanisms of the macrophage and find a safe niche in the phagosome for intracellular growth. $\text{IFN}\gamma$, an effector molecule of the adaptive immune response, halts this unimpeded growth of *Mtb* in the macrophage (60). Although it is clear that $\text{IFN}\gamma$ is highly effective in restricting mycobacterial growth in macrophages (60), the mechanisms through which this is achieved is not fully understood. NOS2-mediated antibacterial pathway is one mechanism that has been extensively studied in the murine model of tuberculosis, wherein it has been demonstrated that $\text{IFN}\gamma$, in conjunction with TNF, upregulates NOS2 and the production of reactive nitrogen intermediates within the phagolysosome, resulting in *Mtb* killing (142). Nitric oxide also reacts with glutathione to form s-nitrosoglutathione that is toxic to *Mtb* (143). Despite this accumulated evidence of $\text{IFN}\gamma$ -mediated antimycobacterial activity in the murine model, the *Mtb* killing mechanism in human macrophages is less clear (144). Individuals with mutations in the $\text{IFN}\gamma$ receptor are more susceptible to mycobacterial infections suggest that $\text{IFN}\gamma$ -mediated antimycobacterial pathways are active in human macrophages (69). Alveolar macrophages of tuberculosis patients express NOS2, and isolation of NOS2-expressing macrophages exhibit antimycobacterial activity *in vitro*, which can be abolished in the presence of NOS2 inhibitors (145). Thus, whether NOS2-mediated pathway is active in human macrophages remains an open question and awaits better methodologies for studying the enzyme.

Recent evidence indicates that $\text{IFN}\gamma$ induced LRG-47, a GTP-binding protein has a principal role in the ability of the host to control *Mtb* replication, since mice lacking LRG-47 are highly susceptible to *Mtb* infection (146). Based on several elegant works, autophagy, originally defined as a cellular homeostatic process, is emerging as a powerful host defense machinery of innate immune cells (147). $\text{IFN}\gamma$, through enhancement of LRG-47 activity, was shown to induce autophagy in *Mtb*-infected macrophages, which resulted in revoking the restriction on *Mtb* phagosome maturation and delivery of the an-

timicrobial contents of the lysosomal compartment (148). Whether autophagy or more specifically “immunophagy,” (a term coined by Deretic to define the specialized function of autophagy in host immunity (147)) restricts *Mtb* growth in human macrophages and whether innate immune signaling pathways such as TLRs induce immunophagy are all very significant questions that need to be addressed.

OTHER INNATE IMMUNE CELLS

Besides macrophages and dendritic cells, $\gamma\delta$ T cells, NK cells, and NKT cells also participate in the innate immune response to TB. Murine studies have indicated that the induction of $\gamma\delta$ T cell in the immune response against TB precedes that of conventional CD4 and CD8 cells and hence plays an important role in modulating the effector response against tuberculosis. Following infection, the early recruitment of cells to the lung is mediated by the chemokines, CXCL2 and CXCL10, and the cytokine, IL-12 released by macrophages and dendritic cells in the lungs (149). Once activated, $\gamma\delta$ T cells secrete $\text{IFN}\gamma$ and $\text{TNF}\alpha$. The production of these cytokine strengthens the bactericidal capacity of macrophages by induction of NOS2. Recently, it has been shown that $\gamma\delta$ T cells secrete IL 17 in response to IL 23 secreted by dendritic cells, thereby implicating them as a main player in the resistance against infection at the initial stage (150). Response of mice deficient in $\gamma\delta$ T cells to *Mtb* infection is dependent on the dose and the route of infection. These knockout mice are able to contain *Mtb* infection with a low inoculum; however, infection with a higher inoculum of *Mtb* administered intravenously resulted in the formation of pyogenic granulomas, indicating that a role for these cells is perhaps in cellular traffic during infection (151, 152). In these experiments with high dose of virulent *Mtb*, the lung pathology indicated enhanced migration of neutrophils and increased size of granuloma thereby indicating a role of $\gamma\delta$ T cells in granuloma formation and mycobacterial containment (152).

The antigen specificity of murine $\gamma\delta$ has not been well studied. However, it has been shown that murine $\gamma\delta$ cells do not respond to phosphate antigens, which are recognized by human $\gamma\delta$ cells. Upon contact with *Mtb*, $\gamma\delta$ cells have been shown to secrete IL-2 and exhibit cytolytic function and hence involved in innate immune effector mechanism (153). In both humans and primates, antigen-specific $\gamma\delta$ cells recognizing phosphoantigens have been documented and have been shown to generate a memory response (154, 155). Loss of $\text{V}\gamma 9 + \text{V}\gamma\delta 2 +$ subset of $\gamma\delta$ T cells was shown to be correlated with tuberculosis (156).

NK cells are recruited to the lungs early during Mtb infection. There they are known to expand and become a primary source of IFN γ . Activated NK cells are known to cause lysis of infected macrophage by utilizing NK cell receptor in TLR-dependent manner (157). NK cell depletion studies have shown no change in bacterial burden (158). However, recent studies (159) indicate that NK cells provide resistance during early Mtb infection via production of IFN γ . The IFN γ activated macrophage in a NOS2-dependent manner and also by regulating neutrophil migration to the lung for controlling lung inflammation. Future experiments defining the exact contribution of NK cells and the IFN γ secreted from them to innate immunity would help understand whether NK cell activation has a role in preventing infection following exposure to Mtb.

NKT cells are TCR-expressing T cells which also express the NK cell marker NK1.1. In mice, NKT cells are mainly represented by V α 14 NKT cells, while in humans, there is a homologous population of V α 24 NKT cells. NKT cells are known to recognize nonpeptide antigens in the context of CD1d. Role of NKT cells in tuberculosis has been studied in both humans and mice. Human V α 24-restricted NKT cells are activated by α -galactosylceramide. CD1d-restricted NKT in the presence of α -galactosylceramide cells restrict the growth of Mtb in a granulysin-dependent manner (160). It has been shown that NKT cells induce a granulomatous response to a glycolipid fraction of Mtb cell wall (161). This finding is further supported by the fact that α -galactosylceramide-activated NKT cells contribute to enhanced resistance against Mtb infection (162).

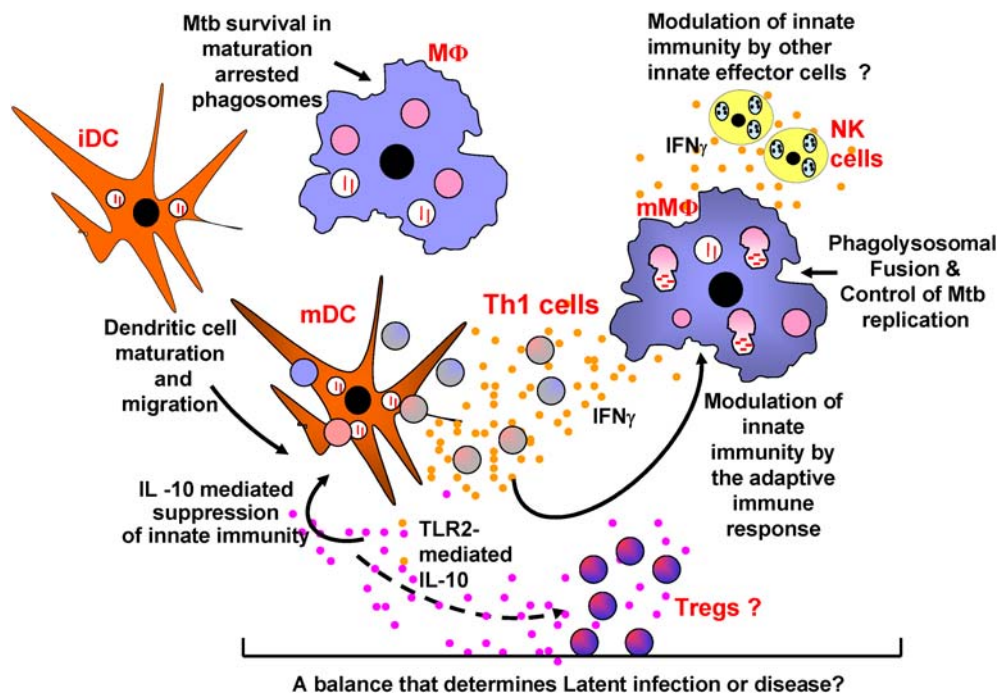


Fig. 2. A paradigm for how the innate immune response to Mtb regulates the adaptive immune response. Initially, Mtb survives and replicates inside macrophages since it can prevent fusion of its phagosome with the lysosomal compartment. Concomitantly, dendritic cells capture Mtb, undergo maturation, and migrate to the draining lymph nodes. Adaptive response is initiated in the draining lymph nodes wherein naïve antigen-specific T cells are primed by dendritic cells to Th1 and cytotoxic effector cell types. Mycobactericidal function of macrophages is dependent on IFN γ , initially produced by innate immune cells such as NK cells and later on provided by effector T cells. The secreted IFN γ promotes phagolysosomal fusion and enhances Mtb killing. TLR2-mediated innate IL-10 is released during the induction of innate immune response and subsequent Th1 induction. The role of the innate IL-10 is to control the magnitude of the Th1 response by either down modulating antigen-presentation function or by inducing T regulatory cells. The TLR2/IL-10 axis, on the one hand, is important for allowing Mtb to achieve the latent state and, on the other hand, may also cause excessive immunosuppression leading to disease. The *unbroken lines* indicate that experimental evidence is available and *dashed lines* indicate that it is speculative and is an area for future investigation.

There is still much to learn regarding the contribution of $\gamma\delta$ T cells, NK cells, and NKT cells to the overall innate immune protection against TB in humans. A next step is to study these cells in the context of protection against clinical strains of Mtb in humans.

CLOSING THOUGHTS

An emerging principle in intracellular parasitism is that successful pathogens such as Mtb have acquired the ability to persist in the host without always inducing disease and mortality (163). The strategy on the part of Mtb is to induce sufficient Th1 immunity in the host to control its replication but not result in its complete eradication. The advantage to the host is minimal collateral damage to lung tissue. Thus, Mtb remains dormant in the host for decades, in a sort of symbiotic relationship. Under certain altered conditions in the host, Mtb will reactivate and cause immunopathology such as lung cavitation, which increases its infectivity and thereby maintains the cycle of transmission to new hosts.

Although Mtb interacts with several different TLRs on host cells, we posit that to establish a latent infection Mtb specifically usurps the innate TLR2 signaling in the host to blunt Th1 immune responses. Supporting evidence for the paradigm include (i) Mtb possess a large gamut of ligands for TLR2; (ii) Mtb/TLR2 interaction suppresses macrophage functions; (iii) innate IL-10 secretion by dendritic cells and macrophages in response to live Mtb is TLR-2 dependent; and (iv) absence of TLR2 results in exaggerated immunopathology in the host. The mechanisms for limiting Th1 response may include inhibition of antigen-presentation functions and induction of T regulatory cells (Fig. 2).

A corollary to the paradigm is that virulent strains can tip the balance toward immunosuppression using the same TLR2 signaling pathway. Although there is no evidence that TLR2 interaction is necessary for protection against Mtb disease, it must be pointed out that these conclusions are drawn from studies performed with laboratory strains of Mtb. It is highly probable that the interaction of clinical strains of Mtb with the TLR2 complex does result in potent immunosuppression in the host. In this regard, it would be interesting to determine if the immunosuppressive cytokines induced by the phenolic glycolipid of the virulent Beijing strains (164) is TLR2-mediated, and whether the induction of T regulatory cells present in TB patients (165, 166) is TLR2-dependent.

Clearly, future studies should investigate if the differing interaction of Mtb clinical strains with the TLR2/IL-10 axis is the control switch that determines whether the

outcome of Mtb exposure is latent infection or tuberculosis disease.

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