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TB vaccines; promoting rapid and durable protection in the lung

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Abstract

TB vaccine discovery has focused on IFN- γ both for the selection of antigens and vaccine delivery strategies. Recent breakthroughs in our understanding of requirement for immunological memory and the expression of immunity to TB in the lung now provides a framework for reconsidering that strategy. We will discuss the status of the TB vaccine field, recent insights into the role of central memory cells and the potential of tissue-resident memory cells in vaccine promoted protection against TB.

Introduction

Tuberculosis (TB) kills around 1.5 million people each year and although novel diagnostic methods and intense treatment schemes like DOTS (directly observed therapy short course) is widely implemented in high prevalence areas, worldwide mortality has not been reduced at the expected or desired rates [1].

Adding to the magnitude and complexity of the TB problem, mortality represents only the tip of the iceberg, as more than 2 billion people worldwide are clinically healthy but latently infected with *Mycobacterium tuberculosis* (Mtb). Latent TB infection (LTBI) provides an infinite source of potential reactivation disease and transmission. These numbers make TB the most widespread infectious agent and together with HIV the top cause of death from infectious diseases [1]. The HIV epidemic is the driving force of the TB epidemic in many countries and results in an increasing number of cases involving MDR-, XDR- and TDR (Multi-, eXtensively-, and Total-Drug Resistant) TB [1].

The Bacillus Calmette-Guérin (BCG) vaccine was developed a century ago and is used extensively in most parts of the world with the exception of Western Europe and North

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America. This vaccine has some protective effect in children but fails to protect against pulmonary tuberculosis in adults. Mtb and humans have co-evolved since the most early human origin [2], and this has allowed the pathogen to adapt and develop a refined set of countermeasures that represent a very difficult target for the immune system.

The result is a pathogen that is rarely cleared by the natural immune response but instead establishes a long-term chronic infection [3]. Attempts to identify adaptive immune responses that are missing or are insufficiently expressed in individuals that develop disease has so far not been successful [4, 5]. However, although clearly insufficient for the control of TB in individuals that develop disease the immune response raised during the natural infection still guides most attempts to develop vaccines.

Vaccine strategies against TB

In the last 10 years, there has been substantial progress in the TB vaccine field, with more than a dozen novel vaccines in clinical trials. These vaccines can be divided into different categories depending on the time point of administration compared to the infection and/or prior BCG vaccination and the delivery system used (see Figure 1 for a schematic representation and description of the vaccines in clinical trials). Live mycobacterial vaccines such as recombinant BCG or attenuated Mtb are intended as replacements for neonatal BCG vaccination.

Preventive booster vaccines aim to prolong and increase the efficacy of neonatal BCG vaccination and are subunit vaccines either based on recombinant antigens in adjuvants or expressed in live viral vectors such as MVA or adenovirus. Post-exposure vaccines are designed to be administered on top of already established LTBI and therefore include antigens that are upregulated by bacteria in this particular stage of the infection. The final strategy is the use of therapeutic vaccines in TB patients to complement or shorten conventional chemotherapy (Fig. 1). For a comprehensive overview of the current status of the different TB vaccines in clinical trials, see e.g. [6, 7].

Most of the novel vaccines are currently in clinical phase 1/2 trials to evaluate safety and immunogenicity. However, recently one of these new vaccines, a modified vaccinia virus expressing Ag85A (MVA85A) was the first new TB vaccine in for more than 60 years to undergo a clinical efficacy trial [8]. During its pre-clinical development program, this vaccine was demonstrated to boost powerful Th1 responses measured as IFN- γ EliSpots, which by most investigators in the field was seen not as a perfect correlate of protection but as the best measure of vaccine take.

The phase 2b clinical trial was conducted in South Africa and BCG-vaccinated infants were boosted with MVA85A and followed for three years. The outcome of the trial was very disappointing with no detectable improvement of protection against TB [8].

As recently pointed out [9], although the BCG-MVA booster strategy was also successful in promoting a very strong Th1 response in animals, it offered no significant increase in protection compared to BCG alone in various animal models. This lack of protection in the face of a strong TH1 response in both the preclinical models and in clinical trials has

emphasized the need to reconsider the immunological requirements for TB vaccines. Our focus will be on the relevance of vaccine responses that are not characterized by high levels of systemic IFN- γ such as central memory and immunity expressed in the lung.

TB: Infection, Immunity and Antigens

Mtb is an airborne pathogen that can establish infection in susceptible individuals when aerosolized water droplets, even those containing very few infectious bacteria, are inhaled. The bacteria are thought to first be taken up by alveolar macrophages and are later acquired by other myeloid cells, including neutrophils and dendritic cells[10]. Importantly, virulent mycobacteria utilize cell surface lipids to restrict TLR-driven recruitment of host-protective immune cells to the initial site of infection [11]. Thus, for the first week after bacteria are inhaled, the host response to Mtb is probably mediated primarily by host cells resident in the lung prior to infection. Furthermore, Mtb manipulates cell death pathways to delay bacterial transport to the lung draining lymph node [12, 13], and upon arrival in the lymph node, Mtb induces an inflammatory milieu that promotes the expansion of highly suppressive pathogen-specific regulatory T cells [14]. These Mtb-driven processes slow the priming of Mtb-specific effector T cells in the lymph node, and delay their ultimate arrival at the site of infection in the lung [15].

The delayed onset of adaptive immunity in the lung is widely believed to facilitate the ability of Mtb to establish a niche for chronic infection [16, 17]. By the time T cells finally arrive at the site of infection they function to contain the bacteria, and play a role in granuloma formation, positioning themselves around infected macrophages (Figure 2). However, rarely are they able to eradicate infection. CD4 T are critically important for TB protection and the cytokines IFN- γ and TNF α are required for control of bacterial growth in both animal models and in man [3, 17]. These cytokines activate macrophages to control bacterial growth by a combination of reactive oxygen and nitrogen intermediates, lysosomal enzyme attack, antimicrobial peptides (such as cathelicidin) and autophagy [3, 17]. Th17 cells can accelerate the initial response and promote the recruitment of other cell types including neutrophils and CXCR5+ CD4 T cells to the site of infection through their secretion of IL17 [18, 19]. CD8 T cell responses increase later during infection often with a kinetic that is associated with increasing bacterial burden. Their role in protective immunity remain unresolved and recent data suggest that even very high numbers of vaccine promoted CD8 T cells specific for protective antigens fail to influence Mtb growth [20, 21].

Mtb has evolved a set of strategies including arrest of phagosome maturation and lysosomal movements, modulation of cell death pathways, prevention of antigen presentation, and release of anti-inflammatory factors that allow it to evade elimination and establish a latent infection [22]. In this stage of the infection, part of the bacterial population transforms into a slow or non-replicative state (so-called dormant Mtb) [23]. LTBI represents a source of continuous antigen exposure that has profound influence on the long-term maintenance of immunity (see below). Mtb can resuscitate whereby the dormant bacteria transforms into a highly replicative and metabolically active stage. The molecular cues that triggers this transformation is still unknown. Alternatively, individuals with LTBI can develop active TB due to airborne infection with a new strain of Mtb, which may occur even more frequently

than reactivation TB in regions of the world in which TB is endemic. It is unknown how ongoing LTBI alters the immune response to new Mtb infection.

The host recognizes multiple antigens during TB infection and the antigens expressed by Mtb changes in different stages of the infection [24, 25]. Therefore, in contrast to the findings for some viral diseases there is not one surface exposed antigen, which is critically important for protection. When comparing the many efforts over the last 25 years to discover novel antigens and in particular recent whole genome screening efforts, it is clear that certain gene families like the ESX gene family are immuno-dominant and very frequently and strongly recognized [26, 27]. Recent data suggest that some of these immuno-dominant antigens contain cryptic epitopes that are not targeted during the natural infection but can still induce a protective immune response [28, 29].

Protective immunity and memory

The goal of all TB vaccines currently in clinical trials is to induce memory T cells capable of mounting a protective response to subsequent Mtb challenge. Animal models suggest that this protective T cell response should have two principal components. First, it should be rapid and robust at the primary site of infection in the lung, and second, it should be durable [17]. Achieving both of these goals through vaccination requires careful consideration of the functional properties of distinct subsets of memory T cells, particularly their propensity to traffic to and to be retained within the lung, both prior to and during ongoing Mtb infection. Mounting a rapid frontline of defense in tissues, including the lung, has long been considered the domain of circulating effector T cells (T_{EM}), but more recently the role of tissue-resident memory cells (T_{RM}) has become increasingly appreciated [30]. Conversely, central memory T cells (T_{CM}) are required to maintain a protective T cell response in the face of chronic antigenic stimulation [31]. Recent advances in our understanding of the advantages and limitations conferred by each of these memory subsets in the context of TB are discussed below.

The limitations of T_{EM}

As the most differentiated of the T_{EM} subsets, T_{EM} of the T helper 1 (Th1) lineage are the most robust producers of $IFN-\gamma$ [31]. Because of this property, the primary goal of most TB vaccine candidates has been to induce a large T_{EM} population. T_{EM} lack expression of the chemokine receptor CCR7, therefore they do not recirculate in the lymphoid system [31]. Rather they reside in the spleen, circulate in the blood, and enter and exit tissue sites, including the lung, as they patrol for foreign invaders. They represent a first line of defense; when tissues become infected, circulating T_{EM} can be recruited quickly to these sites of inflammation. Despite these properties, the limited ability of T_{EM} to prevent TB are becoming increasingly apparent. First, T_{EM} circulating in the blood are poorly recruited to the lung during the early stages of infection [32]. This, at least in part, is likely to occur because of the dampening of TLR-mediated recruitment of immune cells to the lung by cell surface virulence lipids [11]. Later, during the chronic phases of TB, terminally differentiated effector T cells are excluded from the lung parenchyma and accumulate in the lung-associated vasculature [33]. This localization likely severely restricts their protective capacity, as optimal protection is provided only through cognate interactions between Mtb-

specific T cells and infected cells, most of which reside in the lung parenchyma [34]. Finally, T_{EM} are relatively short-lived, and fail to persist in the face of ongoing antigenic stimulation seen during the chronic phase of TB infection [35]. Therefore, vaccines, such as BCG, that prime a large proportion of T_{EM} , confer insufficient levels of long-lived immune protection in animal models [36, 37].

The promise of T_{RM}

In contrast to T_{EM} , T_{RM} comprise a newly defined subset of memory T cells that do not circulate, but take up residence in peripheral tissues, mucosal sites, and barrier surfaces, providing a frontline of defense at these locations [30]. Given the delayed recruitment of T_{EM} from the circulation to the lung after airborne Mtb infection, vaccine-induced lung T_{RM} may theoretically provide the best avenue for early Mtb control (Fig. 2A). Although the rules governing the induction, maintenance, and function of T_{RM} are only beginning to be understood, respiratory viral infections are known to induce T_{RM} that persist in lung tissue and the lung airways following infection clearance [38-40]. Both CD8 and CD4 T_{RM} have been identified, but CD4 T_{RM} seem to be more numerous [41]. Phenotypically, mouse CD4 T_{RM} are distinguished from circulating memory T cell populations by their expression of CD69 and CD11a, whereas CD8 T_{RM} also express CD103 [39, 40]. Importantly, it was recently demonstrated that the high expression of CD69 found on T_{RM} functions to retain the cells in peripheral tissue by interfering with the sphingosine-1-phosphate receptor [42]. In addition to T_{RM} of the Th1 lineage that may provide direct protection against Mtb infection, lung resident Th17 may also play an important role by inducing factors, including chemokines, that expedite the recruitment of Th1 T_{EM} from the circulation into the lung [18, 19]. Given our current understanding, vaccines may need to deliver Mtb antigens directly to the lung mucosa to efficiently elicit lung T_{RM} [41]. The role of T_{RM} in TB protection still needs experimental confirmation but induction of this subset may have the potential to improve early control with infection in preventive vaccine approaches.

The need for T_{CM}

In contrast to T_{EM} and T_{RM} , T_{CM} express CD62L and CCR7 and circulate through lymphoid tissues, but generally have a limited ability to enter infected tissue sites [31]. As a result, protection provided by T_{CM} is delayed and these cells are therefore generally seen as a potent second line of defense. Upon delivery of their cognate antigen to lymphoid tissue, T_{CM} proliferate rapidly and differentiate into multiple effector cell types, including those capable of migrating to tissue sites of infection. Because of their proliferative capacity and their role as pluripotent progenitor cells, T_{CM} are important mediators of long lasting immunity. Chronic TB infection diminish the T_{CM} population by providing a continuous source of antigen that pushes Mtb specific T cell clones to terminal differentiation [36], whereas T cells specific for cryptic epitopes that are not expanded by the ongoing infection may have the ability to bypass this development [29].

The first indication that T_{CM} may have a particularly important role in protective immunity to TB infection in the lung comes from adoptive transfer studies of immune CD4 T cells. The unexpected finding in this early study, was that whereas both CD62L high and low cells could transfer protection to the spleen of recipient mice, protection of the lung was mediated

exclusively by T cells that expressed high levels of CD62L i.e. T_{CM}'s [43]. Recent studies examining the role of T_{CM} in immunity against TB have revealed surprising insights into their trafficking patterns and protective capacity during Mtb infection that allow a rational explanation of this early observation. Despite ongoing stimulation by Mtb antigens, Mtb-specific CD4 T cells with many T_{CM}-like properties are found in the lung during infection [44]. Their maintenance depends on intrinsic expression of ICOS, the transcription factor Bcl-6, and to a lesser extent, CXCR5, all components of signaling pathways operant within lymphoid follicles. The fact that these pathways operate within the lung parenchyma during TB highlights the role of the TB granuloma as a tertiary lymphoid structure within the lung (Fig. 2B) [45, 46].

Most, if not all, of the components that define lymph nodes, including CCL21, CXCL13, follicular dendritic cells, efferent and afferent lymphatics, and high endothelial venules are also present in granulomas [19]. The discovery that, upon adoptive transfer, cells with T_{CM}-like properties [44] migrate directly into the parenchyma of the Mtb-infected lung whereas terminally differentiated effector T cells are excluded [33], was initially surprising and somewhat confusing. However, if movement of T cells into and out of the lung during TB is governed by the rules that dictate trafficking of T cells through granulomatous lymphoid structures; these results begin to make sense (Fig. 2A).

In further support of the idea that lymphoid follicles of the granulomas are critical for directing T cell trafficking and maintaining protective T cell populations, T cells lacking CXCR5, a chemokine receptor expressed by T_{CM} that promotes migration into lymphoid follicles, are limited in their capacity to mediate protection [19]. Taken together, these results help to explain why vaccines that induce large numbers of T_{CM} confer superior and more durable protection compared to other vaccines that induce much greater numbers of IFN- γ producing T cells [36, 47]. Not only do T_{CM} have a high proliferative capacity and the ability for pluripotent differentiation, but also the biology of granulomas as lymphoid structures seems to promote migration of T_{CM} into sites of infection within the lung.

Conclusions

Discovery of new and improved TB vaccines has been complicated and delayed by major gaps in our understanding of how immunity to TB manifests itself at the site of infection in the lung. Without this knowledge, too many efforts in TB vaccine discovery have been dominated by trial and error with the only conclusive read-out being the ability to protect against challenge in experimental animal models. Given that there is no direct correlation between protection and the level of systemic IFN γ promoted by experimental TB vaccines, we now need to develop the next generation of vaccines with specific focus on immunological memory expressed in the lung.

A successful vaccine strategy induce protective pulmonary cellular responses that can rapidly respond to Mtb and it seems logical that T_{RM} localized in the lung parenchyma might be critical in this context. However, the evidence for improved efficacy by aerosol delivery of vaccines is still scarce and restricted to the use of airway virus like adenovirus for vaccine delivery [48]. In preclinical vaccine studies it is very important to keep in mind

that choosing the aerosol route for vaccination may result in misleading interpretations as inflammation and innate immune activation at the port of entry may have both a positive and negative impact on resistance against subsequent challenge [49]. It is therefore critically important to define the role of T_{RM} in TB prevention and in this context to discriminate between short-lived innate activation and long-lived adaptive immune responses. This work will depend on the development of effective approaches to pull the T_{RM} 's into the lung from the peripheral repertoire and to provide the necessary signals for their long-term maintenance at this site.

Given the efficiency with which T_{CM} enter the TB infected lung parenchyma and the longevity of these responses, clearly this subset plays a decisive role in TB vaccines. Eliciting T_{CM} to Mtb antigens that are expressed also during chronic infection may target bacteria localized in lung granulomas resulting in long-term control with infection of particular importance for post-exposure vaccines [50]. Although we still do not fully understand the vaccine parameters that govern the development of T_{CM} 's, several novel adjuvanted Tb subunit vaccines seems to strike a very desirable balance between T_{CM} and T_{EM} in both animal models [36, 51], and in clinical trials [52, 53]. BCG in contrast promotes a response that is much more skewed towards T_{EM} [36, 37] and a recent study demonstrated that increased levels of T_{CM} 's were a prerequisite for the improved efficacy of a novel genetically engineered strain of BCG [54].

One important consequence of this recent improved understanding of immunity to TB in animal models is that to optimize vaccine promoted immune responses in clinical trials we need better tools for monitoring responses in the lung. Typically, TB vaccine clinical trials are designed to assess immune responses in blood samples collected at various time points after vaccination. Recent advances in our technical ability to monitor the expression of multiple cytokines and cell surface receptors have greatly improved our capacity to determine the T_{EM}/T_{CM} ratio in the blood. This, however, only provide limited insight into the local immune responses in the lung, especially those that fully develop only after the establishment of infection.

Even recent attempts to increase the relevance of in vitro assays through biologically relevant read-outs like mycobacterial grow-inhibition [55], may prove futile if the sample is obtained from the blood. It will therefore be critically important, both in preclinical animal models and in clinical trials, to move away from simple measurements of $IFN-\gamma$ in the blood and towards quantifying T_{RM} (prior to infection) and T_{CM} (after infection) in the lung. Studies in animal models will be particularly important to understand the factors that govern the localization of vaccine-elicited immunity to the lung site of infection. Ultimately, hypotheses generated in animal models will need to be tested in human trials, and BAL has proven to be a useful technique for obtaining T cells resident in the lung airways. Due to its invasiveness, this approach may be more feasible in small clinical trials than for larger ones. It therefore remains a top priority to determine if blood assays can provide signatures of efficient vaccine-promoted immunity in the lung. Such assays would provide a big boost for future vaccine discovery, optimization and clinical evaluation.

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- Special interest
 - Outstanding interest
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Highlights

Most TB vaccine candidates aim to amplify IFN- γ producing memory T cells in the blood

The frequency of IFN- γ producing T cells in blood does not correlate with protection

Animal models suggest vaccine-induced T cell responses should be rapid and sustained

Lung resident-memory T cells may prevent or curb early infection

Central memory T cells migrate to the site of infection and provide durable protection

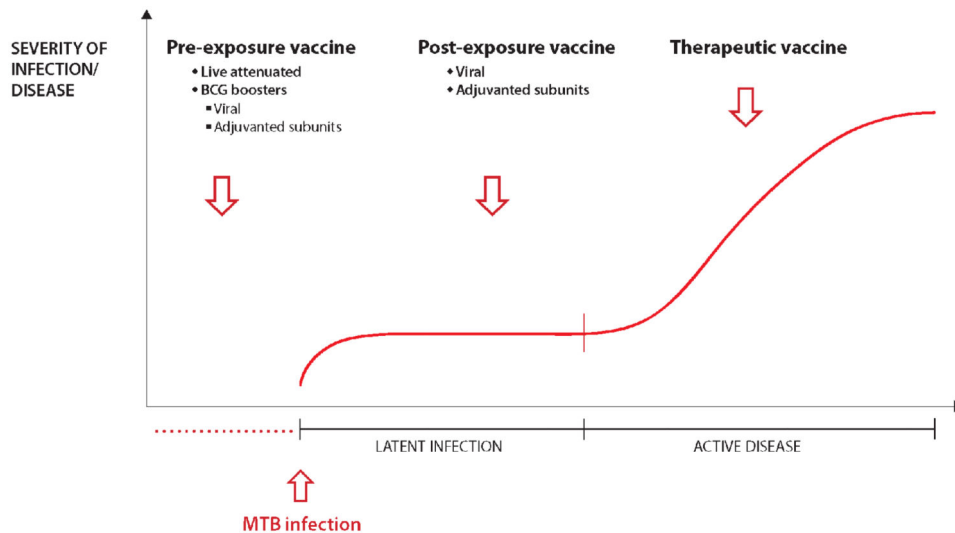


Figure 1. Different types of TB vaccines

Vaccines are divided into different categories depending on the time point of administration and the delivery system used:

Pre-exposure vaccines are administered prior to infection with *Mtb*. Viable mycobacteria designed to replace BCG as prime vaccines are classical pre-exposure vaccines. There are two different vaccines in clinical trials; the recombinant BCG *ureC hly*⁺ (VPM1002) and attenuated mycobacteria with two gene deletions (MTBVAC) [6, 7].

BCG booster vaccines comprise *Mtb* protein antigens expressed in viral vectors or delivered in adjuvant. There are currently several different subunit vaccines in clinical trials based on viral delivery (MVA85A and Crucell Ad35) and protein in adjuvant (H4/IC31 and M72F/AS01E) [6, 7].

Post-exposure vaccines target adolescents and adults with LTBI. The ID93 and H56/IC31 subunit vaccines have been tailored for this strategy by integrating latency antigens of *Mtb* with the goal of enhancing immune pressure and control of infection to prevent reactivation of TB in latently infected individuals [6, 7].

Therapeutic vaccines could be used as an adjunct to conventional chemotherapy to shorten the treatment period and prevent recurrence of disease. The RUTI vaccine is a complex extract of mycobacterial antigens that is currently in clinical trials [6, 7].

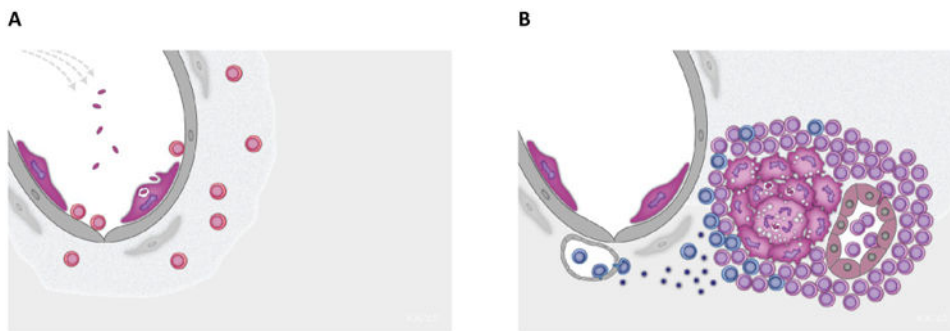


Figure 2. Vaccine promoted immunity in the lung

Vaccine protection against new infection with Mtb (Panel A) or long-term control of Mtb growth in the granuloma (Panel B).

Panel A: T_{RM}'s (in red) localized in the lung parenchyma and airways may be critical for preventing new infections or re-infections. T_{RM}'s are closely related to T_{EM}'s but do not circulate. They reside in peripheral tissues where they mediate a fast and efficient first line defence against invading pathogens. T_{RM}'s can be localized to the lung through aerosol delivery of vaccines.

Panel B: T_{CM}'s (in purple) mediate sustained control with bacterial growth in the granuloma. T_{CM}'s provide a self-renewing source of specific T cells that efficiently migrate into the parenchyma of the Mtb-infected lung. The granuloma acts as an ectopic lymphoid structure which provide the necessary components, such as HEV (to the right - inside the granuloma) to allow T_{CM}'s direct access to the granuloma. T_{EM}'s (in blue) rely on the conventional route into inflamed tissue through small capillaries along a chemokine gradient (to the left – outside the granuloma), but this port of entry seems to operate much less efficiently in the Mtb infected lung.