Protection against tuberculosis: cytokines, T cells, and macrophages

S H E Kaufmann

Tuberculosis remains a major health problem, with two million deaths and eight million new cases annually. At the same time, two billion people (one third of the total world population) are infected with the aetiological agent, Mycobacterium tuberculosis. Of these, fewer than 10% ever develop disease, although the pathogen is not eradicated but rather contained in discrete lesions. Hence, the immune system is highly effective in containing the pathogen, but fails to eradicate it. Disease typically develops through reactivation once the immune system is weakened. The immune response to M tuberculosis is T cell dependent. It comprises not only the conventional CD4 and CD8 T cells, but also γδ T cells and CD1 restricted T cells. γδ T cells recognise phospholipids and no presentation molecules are known thus far. CD1 restricted T cells recognise glycolipids, which are highly abundant components of the mycobacterial cell wall. Although different T cells are required for optimum protection, the immune mechanisms known to have a role in acquired resistance can be associated with two major mechanisms: (a) activation of macrophages by cytokines; (b) direct cytolytic activity. In vivo granuloma formation, which is central to protection, is induced and sustained by cytokines. Mycobacteria are contained within granulomas and in this way are prevented from spreading all over the body.

**Table 1** Tuberculosis: still a major global threat as we have entered the new millennium

<table>
<thead>
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<th>Annual cases of death:</th>
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<tbody>
<tr>
<td>• 56 million cases of premature death</td>
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<tr>
<td>• 17 million cases of death caused by infectious diseases (30%)</td>
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<tr>
<td>• 2.7 million cases of death caused by HIV (4.8%)*</td>
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<tr>
<td>• 2.2 million cases of death caused by tuberculosis (4%)*</td>
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<table>
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<tr>
<th>Annual losses of life years due to disability and disease:</th>
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<tr>
<td>• 1.4 billion total losses of life years</td>
</tr>
<tr>
<td>• 540 million losses caused by infectious diseases</td>
</tr>
<tr>
<td>• 89.8 million losses caused by HIV (6.2%)</td>
</tr>
<tr>
<td>• 33.3 million losses caused by tuberculosis (2.3%)</td>
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</tbody>
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<table>
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<th>Tuberculosis in context:</th>
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<tr>
<td>• 2 billion people infected with M tuberculosis</td>
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<tr>
<td>• 8 million new cases of tuberculosis annually</td>
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<td>• 10–15 people infected annually by a single untreated patient</td>
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*Includes 500 000 people co-infected with M tuberculosis and HIV

**THE DISEASE AND THE UNDERLYING IMMUNE RESPONSE**

Tuberculosis typically is a disease of the lung, which serves both as port of entry and as the major site of disease outbreak. Inhaled droplets containing minute numbers of bacteria are engulfed by alveolar macrophages. These macrophages harbour the pathogen and transport it to draining lymph nodes. A small granulomatous lesion develops containing the bacteria (fig 1). This is true for about 90% of all those who are infected. They will not develop disease directly. Still, because the bacteria are not eradicated, the risk remains that disease may develop later. Although we cannot formally exclude the possibility of spontaneous healing promptly after infection with M tuberculosis, this possibility seems extremely unlikely.

In people with a compromised immune system, however, disease will develop directly after primary infection. This, for example, is the case in patients with HIV infection, in whom the risk of disease development in the following year is dramatically increased depending on the severity of immunodeficiency. Those who contain the infection for years remain at risk and at a later time, when the immune system is weakened, outbreak may occur owing to reactivation. Although considered less likely, reinfection leading to disease is not excluded. Bacterial containment is focused on the granulomatous lesion, where different T cell populations participate in the protective immune response. These include the following T cell populations:

**Abbreviations:** HIV, human immunodeficiency virus; IFNγ, interferon γ; LIF, lymphotixin α; MDR, multidrug resistant; MHC, major histocompatibility complex; TNFα, tumour necrosis factor α

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CD4 T cells recognising antigenic peptides in the context of gene products encoded by the major histocompatibility complex (MHC) class II

- CD8 T cells recognising antigenic peptides in the context of MHC class I
- γδ T cells recognising unusual antigenic ligands independently of specialised presentation molecules—notably, phospholipids
- CD1 restricted T cells recognising glycolipids abundant in the mycobacterial cell walls presented by the CD1 molecules.

MYCOBACTERIA, MACROPHAGES, CYTOKINES, AND T LYMPHOCYTES

*M tuberculosi*s is an intracellular pathogen with the ability to persist in the early phagosomal compartment. It arrests phagosome maturation at an early stage and strongly inhibits phagolysosome fusion. The phagolysosome is a rather hostile environment, where many bacterial pathogens are killed. The early phagosome is a less hostile compartment where *M tuberculosi*s can accommodate itself. Yet, arrest of phagosome maturation by *M tuberculosi*s is not complete and some bacteria are killed or at least prohibited from replication through antibacterial mechanisms including reactive oxygen and nitrogen intermediates, which are produced by activated macrophages (fig 1). The different T cell populations produce interferon γ (IFNγ) and hence are of the T helper 1 (Th1) type. This cytokine is the central mediator of macrophage activation. IFNγ synergises with tumour necrosis factor α (TNFα) in activating macrophages. CD4 T cells also produce lymphotoxin α (LTα), which participates in protection against tuberculosis. At least some of the CD8 T cells, γδ T cells, and CD1 restricted T cells secrete perforin and granulysin which directly kill mycobacteria within macrophages (fig 1).

The costimulatory activity of TNFα in IFNγ activation of anti-mycobacterial macrophage functions was established in the early 1990s. In a series of experiments, Flesch and Kaufmann showed that IFNγ rather than TNFα stimulates anti-mycobacterial macrophage functions in vitro. However, blocking of endogenously produced TNFα reduced IFNγ induced macrophage activation and combination of both TNFα and IFNγ induced optimum macrophage activation. In subsequent studies, the role of TNFα in protection against tuberculosis was elucidated. Gene knockout mice deficient in TNFRI have exacerbated tuberculosis. In particular, granuloma formation in these gene knockout mice is impaired. Moreover, it was found that neutralisation of TNFα in mice with latent tuberculosis results in reactivation and outbreak of disease. These data provide convincing evidence that in vivo TNFα is mostly involved in bacterial containment and in the formation and sustainment of granulomatous lesions. The availability of mice deficient in LTα and LTβ showed that in addition to TNFα, LTα participates in granuloma formation. Hence LTα and TNFα both contribute to granuloma formation, but they do so independently of each other. Figure 2 shows the development of a tuberculous lesion as well as the T cell subsets and cytokines operative in the lesion.

**Figure 1** Different outcomes of infection with *M tuberculosis*, different T cell populations involved in protection, and major anti-mycobacterial effector mechanisms of macrophages. This scheme firstly depicts the different outcomes of tuberculosis in healthy and immunocompromised subjects. Secondly, the figure shows the different T cell populations and their major T cell effector mechanisms in the control of disease. Thirdly, the figure shows anti-mycobacterial effector mechanisms of activated macrophages. (Reproduced from Nature Reviews Immunology (vol 1:20–30). Reprinted by permission from Nature Reviews Immunology (2001;1:20–30). Copyright © 2001 Macmillan Magazines Ltd.)
T LYMPHOCYTE POPULATIONS AS CENTRAL MEDIATORS OF PROTECTION

Because *M. tuberculosis* resides in the phagosomal compartment, its proteins have ready access to the MHC class II processing and presentation machinery. In contrast, it is still incompletely understood how mycobacterial antigens gain access to the MHC class I processing machinery. Similarly, the question of how mycobacterial glycolipids are presented by CD1 molecules remains incompletely understood. CD1b, a major presentation molecule of this pathway, is found in the late phagosomal compartment, which is reached by only very few *M. tuberculosis* organisms (see above). Moreover, macrophages, which are the major habitat of *M. tuberculosis* are virtually devoid of surface expressed CD1 molecules and hence fail to present mycobacterial glycolipids. Some intracellular bacteria, such as *Listeria monocytogenes*, egress from the phagosomal compartment into the cytoplasm. In the cytoplasm, their antigens have ready access to MHC class I processing. It has been proposed that *M. tuberculosis* can also form pores in the phagosomal membrane. Because *M. tuberculosis* apparently remains in the phagosome, it was furthermore assumed that these pores allow exchange of molecules between the phagosome and the cytoplasm. This would not only provide cytoplasmic nutrients to *M. tuberculosis* but also promote antigen translocation to the MHC class I pathway. More recent studies, however, have produced evidence arguing against translocation of phagosomal molecules into the cytoplasm and, as a corollary, against direct antigen transport.

Figure 2: Development of granulomatous lesions in tuberculosis. Promptly after infection, T cells and macrophages are attracted to the site of mycobacterial implantation. There, granulomatous lesions develop. As long as the immune response is competent, the lesions will contain bacteria. These productive granulomas represent a focus of highly dynamic interactions between different T cell populations, macrophages of different maturation stages, and dendritic cells. Once immunity weakens, the balance is tipped and the granuloma can no longer contain mycobacteria. Rather, the granulomatous lesion liquefies and bacteria are released to different tissue sites, different organs, and to the environment. Active disease develops and the patient becomes contagious. (For further details see ref 8.)
tions. Although CD4 T cells, CD8 T cells, and γδ T cells are highly complex comprising different T cell populations, the mechanisms underlying this contain-ment are mostly provided by macrophages. The cross talk between T cells and macrophages is achieved by various cytokines, notably IFNγ, TNFα, and LTα. This cross talk first results in macrophage activation—that is, the ability of macrophages to control mycobacterial replication. Secondly, this cross talk causes the formation of well organised granulomas, where macrophages, dendritic cells, and different T cell populations exist in near vicinity. As long as this cross talk is well balanced, productive granulomas develop which contain the bacterium. Once any element of this complex interplay is impaired, the balance is tipped and the productive granuloma can no longer be sustained. Caseous lesions develop which fail to contain the bacteria. Consequently, bacteria can be disseminated to other tissue sites, to other organs, and even to the environment: active disease develops and the patient becomes contagious. Experiments with gene knockout mice signalling defects have already pointed to a central role for this cytokine in the maintenance of granulomatous lesions and control of disease. Patients with a hereditary defect in TNFα signalling and with tuberculosis have not been described. Therefore, activation of tuberculosis in patients with rheumatoid arthritis treated with anti-TNFα antibodies underlines the importance of this cytokine in the control of latent human tuberculosis.

CONCLUDING REMARKS

Tuberculosis is a chronic disease with protection strongly depending on mycobacterial containment to discrete granulomatous lesions. The mechanisms underlying this containment are highly complex comprising different T cell populations. Although CD4 T cells, CD8 T cells, γδ T cells, and CD1 restricted T cells are apparently all required for optimum protection, a hierarchy seems to exist: CD4 T cells are the most important players, followed by CD8 T cells. The role of these T cell populations in protection against tuberculosis is well understood because of the availability of suitable animal models. The roles of γδ T cells and CD1 restricted T cells are less well understood as appropriate animal models are not available. It is, however, fair to assume that these T cell populations also participate in protective immunity. Although T cells are the major mediators of protection, effector functions are mostly provided by macrophages. The cross talk between T cells and macrophages is achieved by various cytokines, notably IFNγ, TNFα, and LTα. This cross talk first results in macrophage activation—that is, the ability of macrophages to control mycobacterial replication. Secondly, this cross talk causes the formation of well organised granulomas, where macrophages, dendritic cells, and different T cell populations exist in near vicinity. As long as this cross talk is well balanced, productive granulomas develop which contain the bacterium. Once any element of this complex interplay is impaired, the balance is tipped and the productive granuloma can no longer be sustained. Caseous lesions develop which fail to contain the bacteria. Consequently, bacteria can be disseminated to other tissue sites, to other organs, and even to the environment: active disease develops and the patient becomes contagious. Experiments with gene knockout mice signalling defects have already pointed to a central role for this cytokine in the maintenance of granulomatous lesions and control of disease. Patients with a hereditary defect in TNFα signalling and with tuberculosis have not been described. Therefore, activation of tuberculosis in patients with rheumatoid arthritis treated with anti-TNFα antibodies underlines the importance of this cytokine in the control of latent human tuberculosis.

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