

FIRST DR. C.M. SINGH ORATION LECTURE

on

**RECENT ADVANCES ON IMMUNOLOGY
AND IMMUNOPATHOLOGY OF
MYCOBACTERIAL DISEASES**

Delivered at
**Indian Veterinary Research Institute,
Izatnagar (UP)**

on

10th January, 2005

by

Prof. U. Sengupta
MVSc., PhD (Path), FNASC, FAMS
Emeritus Medical Scientist

**Central JALMA Institute for Leprosy
(ICMR)Taj Ganj, Agra 282 001**

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INTRODUCTION

Out of more than 50 known mycobacterial species which comprise the genus *Mycobacterium*, the majority are nonpathogenic environmental bacteria remain in soil and water. A few species which very successfully invade the host and cause pathogenicity are *Mycobacterium tuberculosis*, *M. bovis*, *M. leprae*, *M. ulcerans*, *M. avium* complex including *M. avium* subspecies *paratuberculosis*. As these organisms are very efficient in dampening the defense machinery resident macrophages and dendritic cells of the host by neutralising the antimicrobial properties of cell along with downregulation and upregulation of some of their genes, therefore, these are very efficient in growing inside macrophages. In spite of this host defense builds up its complex machinery of innate and adaptive immunity and makes an effort to compartmentalise the growth of the organisms in immune structures called granulomas. As a result of this mycobacteria generally shuts off its genes to a dormant state which leads to a chronicity of infection. However, in such a situation, most of the time the host may remain asymptomatic with the potential to reappear later in an immunocompromised situation of the host due to other concomitant infection like AIDS. The present review will deal with only the aspects of immunology and immunopathology of mycobacterial infections.

Mycobacterial diseases, chronic granulomatous infections of humans and animals are responsible for worldwide mortality and morbidity of the host. While human tuberculosis is a major health problem causing about 3 million deaths annually in the whole world¹, bovine tuberculosis, is a major cause of economic loss in countries including India where it is endemic and is probably responsible for zoonotic disease problem. *M. tuberculosis* complex consists of *M. tuberculosis*, *M. africanum*, *M. canettii*, *M. bovis* and *M. microti*. While *M. tuberculosis* is a major cause of tuberculosis in humans, in several parts of Africa *M. africanum* is the primary cause of infection². The pathology and course manifested by *M. tuberculosis*, *M. bovis* and *M. africanum* in humans are similar. *M. microti* is a pathogen of voles but avirulent in humans.

M. marinum one of the closest to *M. tuberculosis* complex organism³ is now being used extensively as a model for *M. tuberculosis* pathogenesis⁴⁻⁷.

M. ulcerans, the causative agent of Buruli ulcers, commonly noted in certain areas of the tropics, mostly in West Africa. This organism unlike other mycobacterial species secrete a cytotoxic polyketide toxin which causes extensive, painless, necrotic, noninflammatory ulcers in which bacteria grow extracellularly⁸.

M. leprae, causative agent of leprosy of humans, manifests a wide clinical spectrum between two polar forms, tuberculoid and lepromatous leprosy⁹.

The *M. avium* complex (MAC) is a group related to environmental mycobacteria including *M. avium* subspecies *avium*, *paratuberculosis* and *silvaticum* and *M. intracellulare*^{10,11}.

TUBERCULOSIS

Pathological Manifestations

The epidemiological study and natural history of the disease reveal that most of the hosts are resistant to the infection and can mount a strong immune response to *M. tuberculosis* infection. It is only 10 to 30% of individuals get infected. Of those who get infected 90% or more do not develop disease. In individuals where disease is manifested is mainly due to reactivation of latent infection.

M. tuberculosis is capable of evading innate and adaptive immunity in immunocompetent hosts and evoke sufficient lung pathology for transmission of infection by aerosol for a long period of time. It initially begins with accumulation of macrophages and neutrophils at the site of bacterial lodgment leading to a granuloma formation with epithelioid cells and giant cells around macrophage-laden bacterial growth¹². The granuloma can later undergo caseous necrosis resulting in cavity formation. The host immune defense has a tendency to restrict the growth by laying down a fibrotic layer surrounding the granuloma, which produces microscopic nodules called 'tubercle'. If the host immunity is uphold then this tubercle gets fibrosed and calcified¹²⁻¹⁴.

Interaction between Macrophage and Mycobacteria

Cell mediated immunity (CMI) is primarily expressed by initial macrophage mediated phagocytosis. Although the precise mechanism by which macrophages kill mycobacteria are not clearly understood but several studies have indicated that these cells get activated and gear up their antimicrobial function very efficiently.

Phagosome - lysosome fusion: The lysosomal vacuoles are loaded with active hydrolytic enzymes capable of degrading a wide range of large molecules and microbes. These enzymes function at an optimal acidic pH (4.5-5.0)¹⁵ in the intralysosomal environment. Phagosome containing bacteria after maturation fuses with lysosome forming phagolysosome¹⁶⁻¹⁷. Usually, organisms after an interaction with lysosomal hydrolases undergo extensive degradation within 2 hours of phagocytosis¹⁷. It was shown later that *M. tuberculosis* can prevent phagosome-lysosome fusion resulting in its survival and growth inside macrophages¹⁸⁻¹⁹. It was further noted that viable *M. tuberculosis* containing phagosomes do not fuse and consequently viable mycobacteria are able to grow inside macrophages. Later it was revealed that mycobacterial sulfatides²⁰, derivatives of multiacylated trehalose 2-sulphate, a lysosomotropic polyamionic glycolipid²¹⁻²² are responsible for phagosome-lysosome fusion. Later, in vitro studies showed that *M. tuberculosis* are capable of generation of copious amounts of ammonia in cultures (up to ~20 μ) which has been thought to be inhibitory for phagosome-lysosome fusion²³. Supporting evidence for this also showed that ammonium chloride has inhibitory effect on saltatory movement of lysosomes and has ability to alkalinize intracellular vacuoles²³⁻²⁴. However, finding of the ability of other bases to enhance phagosome-lysosome fusion²³. Contradicted the above finding and therefore the exact mechanism of phagosome-lysosome fusion prevention by ammonia remains to be confirmed again. Mycobacterial enzymes, urease which catalyzes urea to ammonia and carbon dioxide and glutamine synthetase which has a role in alteration of ammonia level through nitrogen metabolism have been under investigation for intracellular survival of *M. tuberculosis*²⁵⁻²⁷.

The RabGTPases are large family of proteins with a variety of regulatory functions in membrane traffic of various endocytic compartments. Mycobacterial phagosomes retain Rab5, which plays a role in interaction between early endocytic compartments and phagosomes and does not allow the phagosome to mature¹⁵⁻¹⁶. It was further noted that *M. tuberculosis* and *M. avium* containing phagosomes can block the vacular ATPase proton pumps and result into reduction in acidification²⁸⁻³⁰. It was later found out that two host proteins, one associated with mycobacterial phagosome; coronin 1/TACO (tryptophan-aspartate-containing coat), an actin binding protein-a component of host phagosome required for phagocytosis and Gal3/Mac2, a protein expressed on the macrophage cell surface and associated with the Golgi and nucleus³³. These proteins are involved in regulating trafficking of molecules in phagosomes. After phagocytosis of mycobacteria these proteins are retained in the phagosome and prevent their maturation and phagosome-lysosome fusion.

Role of Reactive Oxygen Intermediates (ROI) and Reactive Nitrogen Intermediates (RNI): ROI has been shown to be released by activated macrophages induced by cytokines like IFN- γ , TNF- α and these molecules are capable of killing microbes in these macrophages. However, mycobacteria are capable of evading the toxic effect of ROI by several means. Lipoarabinomannan (LAM) and phenolic glycolipid-I (PGL-I), components of *M. tuberculosis* and *M. leprae* are potent scavengers of oxygen radicals³⁴⁻³⁵. Further sulfatides have been shown to interfere with the oxygen mediated antimycobacterial killing by macrophages¹⁴. On the other hand, toxic RNI liberated by activated macrophages have been shown to inhibit the growth of *M. tuberculosis* both in vivo and in vitro specially in mice³⁶⁻⁴⁰.

Role of Toll-like Receptors (TLR): The human TLR is a transmembrane protein with a leucine-rich repeat in extracellular domain of cell. Its cytoplasmic domain is homologous to that of IL-1 receptor. Constitutively active human TLR when transfected in a cell it induces activation of NF- κ B pathway leading to cytokine liberation and co-stimulatory molecule expression⁴¹. The TLR-dependent

activation of the NF- κ B pathway is mediated by an adapter protein, MyD88⁴². For an effective activation and liberation of IFN- γ macrophages require MyD88 adapter protein. MyD88-/- macrophages are five to 100 fold less effective in activating many genes like, nitric oxide synthase, IRG-1, IP10, MIG, RANTES. In vitro studies using *M. tuberculosis* have shown strong evidence that it can signal through TLR2 and TLR4. It has been shown while a soluble heat- stable protease resistant component of *M. tuberculosis* can activate through TLR2, a heat-sensitive cell associated mycobacterial component activate through TLR4⁴³⁻⁴⁴. Recently, it has been noted that prolonged exposure of *M. tuberculosis* LprG (Rv1411c), a 24kDa lipoprotein, induces marked inhibition of MHCII antigen processing. However, LprG was also found to stimulate TLR2 dependent TNF- α production⁴⁵.

Mycobacterial Entry Into Macrophages

In vitro assays using cultured macrophages have shown that the pathogenic mycobacteria can use a variety of receptors to gain entry for their survival and safety inside a cell⁴⁶. It was recently shown that requirement of cholesterol appeared specific for mycobacterial entry⁴⁸, however, whether cholesterol is essential for direct binding with mycobacteria or allowing an appropriate receptor interactions have yet to be determined⁴⁹⁻⁵⁰.

A pivotal role played by macrophage has long been understood in tuberculosis from histopathological and experimental studies⁵¹. In rabbit and mouse models of pulmonary tuberculosis it has been clearly shown that macrophages are responsible for phagocytosis and dissemination of *M. tuberculosis* to the deeper tissues⁵¹⁻⁵³. Recently, infected macrophages with *M. marinum* have been found to migrate from blood stream or hind brain ventricle to deeper tissues in zebrafish. It was also observed that uninfected macrophages from distant areas can engulf dead infected macrophages⁵⁴. Recent comparative gene expression studies using microarray between *M. tuberculosis* log phase cultures and *M. tuberculosis* grown in vitro inside macrophages revealed differential expression of a total of 601 *M. tuberculosis* genes within macrophages relative to that of

axenic cultures. This study further showed that stimulation of infected macrophages with IFN- γ upregulates an additional⁶⁸ activation specific genes in response to generation of NO by macrophages⁵⁵.

Granuloma Formation

After the entry of mycobacterium it is engulfed by macrophages or dendritic cells and are transported to deeper tissues by macrophages and phagocytic cells and additional macrophages start gathering around. Tuberculous granulomas in humans, animals and mice have large component of T lymphocytes, some B lymphocytes, dendritic cells, neutrophils, fibroblasts and extracellular matrix components^{14,56}. CD4+ and CD8+ T lymphocyte subsets play an important role in maintenance of granuloma. The macrophages differentiate into epithelioid cells and in several situations macrophages fuse to become giant cells. A common feature of granuloma is the occurrence of caseation at the centre of granuloma⁵⁵ surrounded by epithelioid cells with a peripheral cuff of lymphocytes (CD4+ and CD8+ T cells)^{56,57}. Mycobacteria mainly reside in the macrophages and in large number in the central caseous zone. In humans and animals the caseous area may become fibrotic and calcified. In a given host at a given time point granulomas can be at different stages of evolution⁵⁸. However, the actual mechanism of caseation still remains unclear. Although, mice infected with *M. tuberculosis* form granulomas without caseation but in rabbits and guineapigs caseations do occur⁵⁹⁻⁶¹.

Immunoregulatory Molecules

Cytokines are responsible for immune functions by the host against the pathogen. *M. tuberculosis* also induces cytokine mediated response for inducing immunity against the infection by the host. The role of the cytokines and their involvement in host response are given as below:

Interleukin-12 (IL-2): *M. tuberculosis* infection induces the host to gear its T helper -1 (Th1) type of immune response. Macrophages and dendritic cells after engulfing *M. tuberculosis* liberates IL-12^{59,61,62} which ultimately leads to IFN- γ liberation and development of Th1 immunity⁶³. It has also been shown that early administration of IL-12 in infected

BALB/c mice significantly lowered bacterial growth⁶⁴. Further, it was shown that IL-12p40 gene deficient mice were susceptible to *M. tuberculosis* infection⁶⁵. A recent study indicated that administration of IL-12 DNA could substantially reduce bacterial burden in mice with chronic *M. tuberculosis* infection⁶⁶.

Interferon-gamma (IFN- γ): IFN- γ is the key cytokine in the Th1 immune response and is essential for the control of *M. tuberculosis* infection in lung^{14,67}. In tuberculosis IFN- γ is produced by both CD4+ and CD8+ Tcells⁶⁸⁻⁷⁰ and by NK cells¹⁴. In the murine system, IFN- γ strongly induces iNOS gene transcript mRNA expression resulting into enzyme activity⁷¹⁻⁷². It has been further noted that IFN- γ knockout mice are most susceptible to virulent *M. tuberculosis* infection⁷³⁻⁷⁴. In human situation where individuals were noted to have defective IFN- γ gene or its receptor were found to be prone serious *M. tuberculosis* infection and other mycobacteria⁷⁵. However, IFN- γ response has been noted in both PPD+ subjects and in active cases of tuberculosis. It has been further noted that the levels of this cytokine is lowered in patients patients of active tuberculosis⁷⁶⁻⁷⁷. In addition it was shown that IFN- γ ultimately is also able to prevent macrophages for production adequate quantum of IFN- γ indicating the limitation of this cytokine in containing *M. tuberculosis* growth in lungs⁷⁸. Recently, it has been shown that aerosol challenge by IFN- γ upregulated only 15 genes of the bronchoalveolar macrophages of tuberculosis patients including IP10 and MCP1⁷⁹. It has already been established that *M. tuberculosis* infection induces IRF-1 (IFN regulating factor1)⁸⁰⁻⁸¹ gene which has pleiotropic function on the immune system⁸² however, the nature of this regulation is still not known. A recent study demonstrated that *M. tuberculosis* inhibits RNA export from the nucleus and affect IRF-1 expression leading to pathogenicity. On the other hand, IFN- γ synergistically stimulates the infected THP-1 cells and upregulates the function of IRF-1⁸³.

Interferon-alpha (IFN- α): It has been reported earlier that IFN- α / β system helps host defense against *M. tuberculosis* infection in mice⁸⁴ and the infection induces human

macrophages and dendritic cells for the production of IFN- α/β ⁸⁵⁻⁸⁶. Further aerosolized IFN- α in combination with chemotherapy has been shown to improve the clinical condition of tuberculosis⁸⁷. Later it was noted in vitro culture system that IFN- α/β released by *M. tuberculosis* infected macrophages inhibit the tyrosine phosphorylation of signal transducer, STAT-1 indicating that it is cellular signal transduction and associated with pathogenicity⁸⁸. Therefore, further research is needed to find out the exact role of IFN- α in immunomodulation of tuberculosis.

Interleukin-4 (IL-4): IL-4, inducer of Th2 response, has not been found at a very high level in tuberculosis patients by most of the authors⁷⁶⁻⁷⁷ however, it has been found to induce apoptosis of *M. tuberculosis* reactive T cells in presence of TNF- α ⁸⁹. *In situ* demonstration of elevated levels of IFN- α expression in granuloma of lymph nodes of patients with pulmonary tuberculosis and tubercular lymphadenitis have been noted with low levels of IL-4 mRNA expression⁷⁶. In experimental situation in gene knockout or ILp40-/- mice no shift to Th2 was observed^{74,65}. Even when granulomas were investigated for IL-4 expression by *in situ* hybridization in advance tuberculosis 3 out of 5 patients showed presence of IL-4 but never in absence of IFN- γ expression⁹⁰. From the above it is clear that IL-4 levels do not correlate with clinical status of tuberculosis.

Tumour Necrosis Factor-alpha (TNF- α): The cytokine, TNF- α plays important roles in both immunity and pathogenesis of tuberculosis. *M. tuberculosis* is known to induce macrophages, dendritic cells and T cells for secretion of TNF- α ⁹¹⁻⁹⁴ which control *M. tuberculosis* infection. Mice deficient of TNF- α or TNF- α receptor are more susceptible to *M. tuberculosis* infection than normal mice⁹⁵⁻⁹⁶. TNF- α along with IFN- γ is known to induce expression of NOS2^{72,90,97}. Further sufficient data have been accumulated to establish the importance of this cytokine in granuloma formation in tuberculosis and other mycobacterial diseases⁹⁶⁻¹⁰⁰. In murine model deficiency of TNF- α leads to a reduction in granuloma formation with fewer epithelioid cells⁹⁹ and impaired

colocalization between lymphocytes and macrophages⁹⁶. However, several mechanisms of TNF- α which promotes granuloma formation are yet to be determined.

The other major function of TNF- α that it destroys the host tissue and leads to pathological progression of the disease in the lung¹⁰¹⁻¹⁰³. Although TNF- α is responsible for tissue damage, it was noted that thalidomide treatment downregulated the cytokines TNF- α , IL-6, IL-10 resulting in a reduction in the size of the granuloma but did not bring about any change in the bacterial number¹⁰³. However, it was later shown in mouse model that neutralization of TNF- α resulted in 100% mortality and the lungs of the mice showed extreme involvement with disorganized granulomas, diffuse cellular infiltration and pathologic manifestations of chronic inflammation¹⁴. These observations rather suggested that TNF- α might play an important role in limiting the pathologic response in tuberculosis. Considering the above it is clear that further study is required for understanding the role of TNF- α in tuberculosis.

Interleukin-10 (IL-10): IL-10 is known as an antiinflammatory molecule which is known to neutralize the function of proinflammatory cytokines. This cytokine during *M. tuberculosis* infection is produced by the macrophages and T cells of the host and downregulates IL-12 production resulting in decrease in IFN- γ level. IL-10 has been found to directly inhibit the CD4+T cell responses, as well as APC function of cells infected with mycobacteria¹⁰⁴. Transgenic mice, constitutively expressing IL-10 were unable to clear BCG infection inspite of normal T cell responses with IFN- γ production¹⁰⁵. Lastly, IL-10 $^{-/-}$ mice were not found to be more resistant to acute *M. tuberculosis* infection as compared to wild type mice¹⁰⁶. Further, the role of IL-10 involvement in pathology and protection are being worked out.

Interleukin-6 (IL-6): IL-6 is known to modulate immune response. It has been shown to suppress T cell responses to antigens in BCG infected macrophages¹⁰⁷. In IL-6 $^{-/-}$ mice early increases with decreased IFN- γ level compared to control mice following a low dose of aerosol infection has been noted¹⁰⁸. Although, these mice with low dose of infection survived but

with high dose of *M. tuberculosis* IL-6^{-/-} mice were susceptible to infection.

Transforming Growth Factor- β (TGF- β): TGF- β is an antiinflammatory cytokine and has been responsible for suppression of T cell responses in tuberculosis patients¹⁰⁹. *M. tuberculosis*¹¹⁰ and LAM¹¹¹ induce human monocyte to produce TGF- β which has been found in granulomatous lesions. This interleukin may also inhibit macrophage in producing NSO2 by IFN- γ ¹¹¹. Increased level of TGF- β 1 have been observed in the pleural fluid and plasma of tuberculous pleurisy patients¹¹²⁻¹¹³. TGF- β has been used to induce pleurodesis in animal models and appeared to be a good candidate for use in humans¹¹⁴. However, an increased level of TGF- β have been associated with increase severity of tuberculosis. PBMC from patients with advanced pulmonary tuberculosis produced significantly higher levels of TGF- β in the absence or presence of whole sonicated antigen of *M. tuberculosis* than PBMC from patients with mild to moderate disease and PBMC from healthy control patients¹¹⁵.

T Cell functions during *M. tuberculosis* infection

Status of CD4⁺ T cells: Experimental models in mice established that CD4⁺ T cells are primarily responsible for protection against *M. tuberculosis* infection¹¹⁶. Anti-CD4⁺ antibody depletion of CD4⁺ T cells¹¹⁷, adoptive transfer^{116,118} or the use of CD4 gene deleted mice¹¹⁹⁻¹²⁰ strongly established that CD4⁺ T cells are essential for controlling infection. In humans important role of CD4⁺ T cells established from the cases of reactivation of tuberculosis in HIV infected individuals in which CD4⁺ cells undergone reduction¹²¹. Further, in HIV-infected patients with tuberculosis, mycobacterial load increases as CD4 depletion becomes more prominent¹²². In addition, a paucibacillary *M. tuberculosis* infection, tuberculous pleuritis, a self-resolving disease, is associated with proliferation of CD4⁺ T cells with high levels of IFN- γ ¹²³. The mechanism by which CD4⁺ T cells contribute to protection against *M. tuberculosis* is due to induction of Th1 cytokines, IL-2 and IFN- γ ¹²⁴ which are sufficient to activate macrophages for intracellular killing of the organism. In MHC II^{-/-} or CD4^{-/-} mice, levels of IFN- γ is severely lowered

during the early stage of infection. Further experiments in mouse model indicated that when with a low dose *M. tuberculosis* infection a chronic infection is generated, treatment with anti-CD4 antibody can evoke reactivation of the infection leading to a high bacterial burden in the lungs causing death of mice¹²⁵. However, IFN- γ levels of CD4-depleted and control mice were same due to IFN- γ production by CD8+ T cells. Further, there was no change in the NOS2 production or activity in the CD4+ T cell depleted mice indicating that there are IFN- γ and NOS2 independent CD4+ T cell dependent mechanisms for control of infection.

Priming of CD4+ T cells, a major function of dendritic cells *in vivo*, is less likely to be affected by *M. tuberculosis* because it was noted that there was no reduction in the levels of MHCII on the cell surface after *M. tuberculosis* phagocytosis^{92,96}. This is further supported by the fact that a strong CD4+ T cell mediated immune response could be generated during *M. tuberculosis* infection. However, a defective T cell stimulation may be contributed by by APCs due to production of cytokines like TGF- β , IL-6 or IL-10 as mentioned earlier. It is very clear that *M. tuberculosis* is equipped with immune evasion strategies by underexpression and overexpression of certain genes when it is engulfed by macrophages or dendritic cells⁵⁵.

Status of CD8+T cells: Experiments in mice definitely proved the protective role of CD8+ T cells¹²⁷. MHC class I presentation of several cytoplasmic antigens is very efficient in tuberculosis. Mice genetically deficient in β 2 microglobulin or TAP in CD8+ T cells were quite susceptible to *M. tuberculosis* infection¹²⁷⁻¹²⁹. Recent studies in mice demonstrated migration of CD8+ T cells to the lungs like CD4+ T cells following *M. tuberculosis* infection^{93,130}. These cells are able to produce IFN- γ and lead to lysis of infected macrophages^{93,127,130}. These interactions between CD8+ T cells and infected macrophages may be manifested in class I restricted or non class I restricted fashion.

Class I restricted CD8+ T cells: In humans and mice, several antigens, for example, 38kDa 131, 65kDa-heat shock protein 132 and 19kDa¹³³ have been shown to be recognized by CD8+

T cells in MHC class I restricted manner. These T cells have been further shown to secrete IFN- γ with lysis of infected macrophages¹⁴. Although the exact mechanism of gaining entry into the cytoplasm of APCs is not clearly understood, however after entry of mycobacteria in the macrophages, the bacilli have been noted outside the phagosome after 4 to 5 days of phagocytosis¹³⁴ and the presentation of the antigen by these infected macrophages can begin at 12 hr which can be later recognized by T cells¹³⁵. It has been shown that live but not heat-killed *M. tuberculosis* infection of macrophages facilitated MHC class I presentation of soluble ovalbumin in a TAP-dependent manner indicating that ovalbumin taken up by phagosomes along with *M. tuberculosis* leaked into the cytoplasm¹³⁶. Further, in BCG infected macrophages proteins up to 70kDa could be noted in the cytoplasm¹³⁷.

Nonclassically restricted CD8+ T cells: CD1 molecules are nonpolymorphic antigen presenting molecules which have been grouped as Group I (CD1a, b and c) and as Group II (CD1d), CD1d molecules like MHC1a molecules, present lipids/glycolipids to T cells¹³⁸. CD1 restricted T cells may be either CD4-8-, CD8+ or CD4+ T cells¹³⁹. Several mycobacterial antigens, such as, mycolic acid¹⁴⁰, LAM¹⁴¹, phosphatidyl inositol manoside¹⁴², glucose monomycolate¹⁴³ and isoprenoid glycolipids¹⁴⁴ have been shown to be presented to the T cells by CD1. Recent study shows that mycobacterial lipids within the phagosomes can be transported in and exported from the cell to endocytic vesicles and can be taken up by any bystander cells for presentation to T cells¹⁴⁵. Although the exact role of CD1 restricted cells in innate and adaptive immunity is yet to be found out, initial studies indicate that CD1 restricted response could be generalized from PBMC of both PPD+ and PPD-subjects^{140,146}. Another study using isoprenoid glycolipid as an antigen could note PBMC proliferation from PPD+ and not PPD-subjects¹⁴⁰. This proliferation further could be blocked by anti-CD1c antibody¹⁴³. CD1 molecules are found usually on dendritic cells with low level of expression on macrophages¹⁴⁷. It is possible that dendritic cells which are in majority in lungs might be involved in stimulating T cells in CD1 restricted

manner in the granuloma. Dendritic cells can be infected with *M. tuberculosis* and it is possible that these cells are playing a pivotal role in protection against tuberculosis. Therefore, research using dendritic cells are needed to unravell various mechanisms which are involved in protection against tuberculosis.

Effector function of CD8+ T cells and protective immunity:

The effector function of CD8+ T cells are primarily expressed by production of IFN- γ and cytotoxicity of the infected cells. Whether both of these functions are conferring protection in tuberculosis is largely unknown. However, it is known that specifically sensitized CD8+ T cells liberate IFN- γ upon TCR engagement or by interaction with *M. tuberculosis* infected dendritic cells in mice⁹³. However, the quantum of IFN- γ produced in such a situation was much lower than CD4+ T cells. Therefore, CD8+ T cell interaction with sensitized macrophages/dendritic cells needs further study. The function of CD8+ T cells in causing lysis of infected cells via perforin and granzymes or the Fas/FasL pathway. Reduction in intracellular mycobacterial number was noted by observing lysis of infected human dendritic cells and macrophages by CD4- MHC classI restricted CD8+T cells¹⁴⁸⁻¹⁴⁹. It was further noted that the perforin was responsible for killing of intracellular mycobacteria¹⁴⁸. In another study it was noted that perforin was responsible for forming pores and the cytotoxic molecule, granulysin was required for lysis of mycobacteria¹⁴⁹.

Status of $\gamma\delta$ T cells: Increasing evidence is being accumulated regarding the role of $\gamma\delta$ T cells in immunity against tuberculosis¹⁵⁰.

These cells recognize a wide range of antigens including the unprocessed protein antigens¹⁵¹. Although studies in mice already indicated the important role of $\gamma\delta$ T cells in early protection¹⁵²⁻¹⁵⁴ such observations in humans and animals have not yet been established. However, higher frequency of antigen-specific $\gamma\delta$ T cells especially bearing V γ 9 V δ 2 cells¹⁵⁵⁻¹⁵⁶ having their ability to produce IFN- γ and cytotoxicity have been recognized in protection against tuberculosis¹⁵⁷.

BOVINE TUBERCULOSIS

It is predominantly caused by *M. bovis*, is a major cause of economic loss in countries where it is endemic and in some countries it has a significant zoonotic problem¹⁵⁸.

The mechanism of pathogenesis and granuloma formation is similar to *M. tuberculosis* in man which has already been detailed above.

It was noted by several workers that IFN- γ response is the major interleukin response in infected cattle¹⁵⁹⁻¹⁶². When both IFN- γ and IL-4 responses to the antigenic challenge (ESAT6) were monitored then it was noted that the IFN- γ level was at a high level all throughout the disease process. On the other hand IL-4 response was delayed and on the 8th week after challenge there was a peak response which declined later. However, both IFN- γ and IL-4 levels were able to discriminate between infected and noninfected cattle in the field in the developed countries¹⁶³. Utilizing these interleukin responses no data are available from the underdeveloped and developing countries.

JOHNE'S DISEASE

Johne's disease, also called paratuberculosis, is a chronic granulomatous enteritis of ruminant animals caused by *M. avium* subsp. *paratuberculosis*. Although Johne's disease can cause death of the animal however, due to the chronicity of infection the morbidity of animal is much more and therefore the disease lay significant impact on the economy of the country¹⁶⁴⁻¹⁶⁵.

Disease Progression and Immunopathology

The disease typically progresses through three distinct stages¹⁶⁶. First and second stages are called subclinical stages where animals are generally asymptomatic. In the second stage also animals remain asymptomatic without developing clinical disease or might progress to clinical stage after 3 to 4 years of infection. In the first subclinical stage the animal is infected but shedding of bacilli in the feces cannot be detected. In the second subclinical stage, the intermittent excretory phase, the bacilli may be discharged in the feces intermittently and not continuously. The third stage is the clinical excretory phase which is characterized by persistent shedding of high numbers

of bacteria in the feces with severe diarrhoea associated with weight loss.

M. avium subsp. *paratuberculosis* enters tissue through M cells present in the distal ileum of calves and goats¹⁶⁷⁻¹⁶⁸. In the Peyer's patches in the intestinal tracts and the jejunum of ruminants adaptive immune response is initiated with T and B cell responses¹⁶⁹.

Fibronectin attachment proteins (FAPs) are family of fibronectin (FN)-binding proteins present in several species of mycobacteria¹⁷⁰⁻¹⁷⁴. Attachment and internalization of *M. bovis* BCG, *M. leprae* and *M. avium* subsp. *paratuberculosis* by epithelial cells *in vitro* has been shown to be dependent on the interaction between FAP and FN^{172,175,176}. B1 integrins have been identified as the host cell receptor for FN-opsionized mycobacteria *in vitro*¹⁷⁴⁻¹⁷⁵. M cells are unique which express high density B1 integrins on the luminal surface¹⁷⁷ and FN bound by organisms simply explains the portal of entry through M cells. Recently, using gut loop assay technique in mice the specificity of M cell-entry through formation of FN-bridge formed between FAP-of *M. avium* subsp. *paratuberculosis* and integrins of M cells have been demonstrated¹⁷⁸.

Lesions associated with the disease is often diffuse and granulomatous and are typically restricted in the ileum and particularly to the ileocecal valve region of small intestine¹⁷⁹. The pathology is mainly due to severe immune pathology induced by immune cells and chronic inflammation¹⁸⁰⁻¹⁸¹. Draining lymph nodes near the lesion often exhibit hyperplasia with large numbers of T cells and macrophages^{180,182} in infected ileal tissues and in associated lymph nodes.

After the initial exposure to the organism, a proper T cell response characterized by the release of proinflammatory cytokines like IFN- γ , IL-1a, IL-6 along with production of IL-2 is generated by the host¹⁸³⁻¹⁸⁴. However, in most of the cases the disease progresses and the Th1 immune response gets shifted to Th2 type of response with IgG1 type of antibody response against the organism¹⁸⁴⁻¹⁸⁶. A recent study using cytokine gene expression in PBMC and tissues of infected cattle indicated that only in subclinically infected cattle IL-10 gene expression was consistently

enhanced in PBMC after stimulation with *M. avium* subsp. *paratuberculosis*. In ileal tissues of infected cattle, expression of genes encoding IFN- γ , TGF- β , IL-5 and IL-8 was greater than the expression in comparable tissues from control uninfected cattle. However, expression of gene encoding IL-16 was lower in tissues of infected than in control tissues. The draining mesenteric lymph node tissues from the infected site showed higher expression of TNF- α , IL-8, IL-2 and IL-10 genes than similar tissues from control animal. In contrast, genes encoding TGF- β and IL-16 were expressed at lower levels in lymph nodes from the infected cattle than the uninfected cattle¹⁸⁷. The above study suggested that cells or other mechanisms capable of limiting proinflammatory responses to *M. avium* subsp. *paratuberculosis* develop in Johne's disease to control tissue damage and immune responses.

LEPROSY

Leprosy, caused by *M. leprae*, is a chronic infectious disease inflicting human beings since the ancient times. As the world by now has attained the targetted prevalence of less than 1 per 10,000 population, therefore, the disease is at present is considered as a disease of no consequence. In India, the prevalence is still less than 3 per 10,000 and it is expected that by another few years the prevalence will come down to less than 1 per 10,000. However, in India still in certain states like Bihar, Uttar Pradesh, Madhya Pradesh, Orissa and West Bengal the prevalence rate is very high and inspite of multidrug therapy the new case detection rate is not going down indicating the existance of man to man transmission of the disease.

The Disease

The disease is expressed with a variety of manifestations depending on the immunity of the host. It may involve nerves (mainly peripheral nerves), skin and mucous membrane. The disease sometimes remain restricted to nerve only and then it is called as pure neuritic leprosy. Considering all the parameter like clinical, bacteriological, histological and immunological the disease has been classified in a spectrum of five-group system with two polar ends. In one end of the pole is

tuberculoid (TT) type and in the other end of the pole is lepromatous type(LL) of leprosy. In between these poles there are borderline tuberculoid (BT), borderline (BB), borderline lepromatous (BL) forms⁹.

It has been noted that at the TT/BT end the level of CMI towards *M. leprae* remains highest and as the disease progresses downwards in the spectrum towards LL pole the CMI towards *M. leprae* goes down as determined by lepromin skin testing and *M. leprae* specific lymphocyte proliferation assay. On the other hand, the level of anti-*M. leprae* antibody and bacteriological load go higher and higher with the advancement of the disease. In this disease there exists a clear inverse correlation between CMI and humoral immunity as well as bacteriological load¹⁸⁸.

Characteristics of T- Lymphocytes in the Granuloma

Extensive studies of T lymphocyte populations have been performed by researchers using monoclonal antibodies for immunophenotyping of cells in the lesions. It was noted that while proportion of CD4+/ CD8+ ratio was more in TT/BT lesions in BL/LL leprosy the ratio was reversed¹⁸⁹⁻¹⁹¹. It has been recently observed that although the CD4+/CD8+ ratio of 2:1 in blood and lesion is not altered in TT leprosy, T-memory (CD4+CD29+) and T-naïve (CD4+CD45+) ratio is 1:1 in blood and 14:1 in lesions¹⁹².

In-situ Environment in Granuloma

The pattern of distribution and localization of various immune cells in the granuloma and epidermal layer of skin help in understanding of the immunological microenvironment. In tuberculoid leprosy the CD4+ T cells remain scattered inside the granuloma along with macrophages, while CD8+ T cells remain in the mantle of the granuloma in the form of a ring^{190,193,194}. It has been established that the above pattern of cell arrangement is an indicator of an immune granuloma. In tuberculoid leprosy, the presence of CD4+ cells, memory T cells and macrophages point towards an active immune interaction with activation and maturation of immune cells, leading ultimately to destruction of *M.leprae*. On the contrary, finding of predominantly CD8+ cells, macrophages and a few CD4+ cells

inside the granuloma indicate generation of a suppressive function in lepromatous leprosy. In addition to the above, the finding of more numbers of CD1+ Langerhan's cells in epidermis and the periphery of tuberculoid granulomas than in lepromatous lesions suggests an environment of effective immunity in tuberculoid leprosy (reviewed in 191).

Th1/Th2 functions of lesional cells: Spectral manifestations in human leprosy have provided the immunopathologists a great scope in investigating the cytokine secretion profile to unravel a variety of immunological responses to the intracellular parasite, *M. leprae*. It has been clearly demonstrated that the majority of *M. leprae* responsive T cells generated from PBMC or skin lesions of TT patients produce IFN- γ , IL-2 and TNF- α but little or no IL-4, IL-5 and IL-6. Further, using immunoperoxidase techniques IL-2, IFN- γ and TNF- α positive cells have been shown to be present in higher number in tuberculoid leprosy compared to that of lepromatous leprosy. Using *in situ* hybridization and mRNA-based PCR further proof for the presence of IFN- α , IL-2, TNF- α , LT and GM-CSF-producing T cells in TT leprosy has been documented. Contrary to the above, these interleukins were found to be at lower levels in BL/LL lesions. However, higher levels of mRNA for IL-4, IL-5 and IL-10 were present in BL/LL than in TT/BT lesions (reviewed in 191). This differential compartmentalization between Th1 and Th2 are almost similar to the dichotomy of Th1 and Th2 which exists in strains of mice¹⁹⁵. Further, functional analysis of T cell clones from lesions revealed that CD4+ clones from tuberculoid leprosy liberating mainly IFN- γ and CD8+ clones from lepromatous lesions produced IL-4¹⁹⁶. It was subsequently noted that some CD8+ clones from LL produced high levels of IFN- γ in addition to IL-4. It was also observed that IL-4 and IFN- γ coproducing clones gradually shifted to Th2 type, whereas non-IL-4 producing clones secreted very high levels of IFN- γ on prolonged cultures¹⁹⁷. In addition when lesions from BL patients were analyzed, it was noted that Th1 type of cells produced IFN- γ and TNF- α with minimal IL-6 production, whereas Th2 type of cells produced IL-4, IL-5, IL-13, IL-10 and Th0 cells produced both types of cytokines¹⁹⁸.

Simultaneous measurement of memory T cell 1 (MT1, CD45RA-, CD62L-, CD11abright, IFN- γ biased) and memory T cell 2 (MT2, CD45RA-, CD62L+, CD11adim; IL-4 biased) of peripheral blood of leprosy patients showed that the ratios of MT1/MT2 differed significantly between tuberculoid and lepromatous leprosy patients¹⁹⁹.

Status of $\gamma\delta$ and $\alpha\beta$ T cells

It has been observed that mycobacterial exposure leads to expansion of $\gamma\delta$ T cells and $\gamma\delta$ T cells have been found in the skin lesions of leprosy²⁰⁰. It was also noted that 25 to 30% of CD3+ T cells of both lepromin skin reaction and reversion reaction are $\gamma\delta$ T cells. The finding of proliferation of $\gamma\delta$ T cell lines in expanded culture generated from skin lesions and peripheral blood in response to *M. leprae* suggests their role along with cytokines in granuloma formation²⁰⁰.

CONCLUSION

Mycobacterial diseases are chronic diseases and mostly remain under control in a host expressing adequate immunity. It is only when there is lowering in the level of CMI (TH 1 type), the focus of containment of infection gets reactivated and the disease is clinically manifested. As the disease progresses there is a shift of Th1 immunity to Th2 type of immunity. However, in tuberculosis no clear cut transformation from TH1 to Th2 type of immune response with progression of disease has been noted as has been seen in leprosy or Johne's disease. Therefore this complex immunological facets have to be understood in a greater detail in future. Further, the mechanism of early events of host-pathogen interaction largely remains unknown. To unravel this, local immunity has to be understood at a greater detail. It is expected that using all the modern techniques in immunobiology a clear picture of the immunological network against the pathogen would be brought out in the near future which might help in designing a suitable vaccine for protection against mycobacterial diseases.

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