

Non-Opsonic Recognition of *Mycobacterium tuberculosis* by Phagocytes

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Key Words

Innate immune receptors · Non-opsonic receptors · Mycobacteria

Abstract

The interactions between *Mycobacterium tuberculosis* and host phagocytes such as macrophages and dendritic cells are central to both immunity and pathogenesis. Many receptors have been implicated in recognition and binding of *M. tuberculosis* such as the mannose receptor, dendritic-cell-specific intercellular adhesion molecule-3 grabbing nonintegrin, dectin-1 and complement receptor 3 as well as Toll-like receptors, scavenger receptors and CD14. While in vitro studies have demonstrated clear roles for particular receptor(s), in vivo work in receptor-deficient animals often revealed only a minor, or no role, in infection with *M. tuberculosis*. The initial encounter of phagocytic cells with mycobacteria appears to be complex and depends on various parameters. It seems likely that infection with *M. tuberculosis* does not occur via a single receptor-mediated pathway. Rather, multiple receptors play different roles in *M. tuberculosis* infection, and the overall effect depends on the expression and availability of a particular receptor on a particular cell type and its triggered downstream responses. Moreover, the role of membrane cholesterol for *M. tuberculosis* inter-

actions with phagocytes adds to the complexity of mycobacterial recognition and response. This review summarizes current knowledge on non-opsonic receptors involved in binding of mycobacteria and discusses the contribution of individual receptors to the recognition process.

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Introduction

Tuberculosis is the world's leading bacterial cause of death, and about one third of the world's population is now infected with *Mycobacterium tuberculosis*. As a facultative intracellular pathogen, *M. tuberculosis* enters the host typically via aerosols, and alveolar resident macrophages are considered the first cells to engulf *M. tuberculosis* and become infected. After this first encounter, dendritic cells and monocyte-derived macrophages also participate in the phagocytic process [1]. Mycobacterial binding to macrophages occurs in cholesterol-rich domains of the host cell plasma membrane [2], and phagocytosis involves different kinds of receptors which either bind to non-opsonized *M. tuberculosis* or recognize opsonins on the mycobacterial surface. These receptors include C-type lectins such as the mannose receptor (MR), dendritic-cell-specific intercellular adhesion molecule-3

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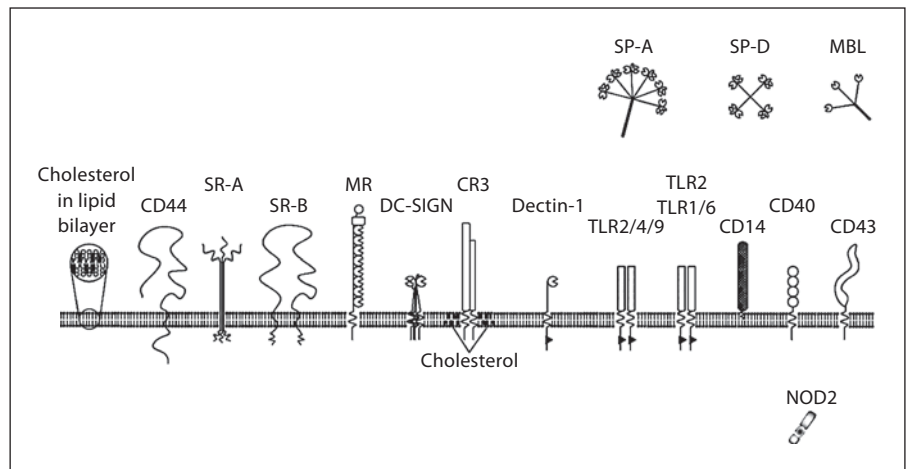
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Fig. 1. Receptors proposed to play a role in recognition and/or uptake of *M. tuberculosis* (illustration by I. Jastram).



grabbing nonintegrin (DC-SIGN) and dectin-1, complement receptors, surfactant protein receptors, scavenger receptors, and glycosylphosphatidylinositol (GPI)-anchored receptors such as CD14 [3, 4]. Other surface molecules such as Toll-like receptors (TLRs), are also important for mycobacterial interactions with phagocytic cells and their subsequent activation [5]. Invasion of macrophages by the tubercle bacillus, which is a facultative intracellular pathogen, is critical in the establishment of infection; consequently, *M. tuberculosis* has evolved several survival strategies and even growth inside its niche in the host cell, the mycobacterial phagosome [6]. *M. tuberculosis* interferes with the Rab-controlled membrane trafficking and thereby prevents phagosome maturation, which involves the fusion of phagosomes with lysosomes. This process, phagosomal arrest, is critical for *M. tuberculosis* persistence as it occurs at a stage where no harm can be done to the bacterium while delivery of nutrients continues [reviewed in ref. 6]. It has recently been demonstrated that tryptophan-aspartate-containing coat protein (TACO), also known as coronin 1 or P57, is indispensable for survival of mycobacteria in phagosomes [7]. This protein is normally released from phagosomes prior phagosome fusion with, or maturation into, lysosomes, but mycobacteria can actively retain TACO on phagosomes, thereby preventing their delivery to lysosomes [8]. TACO was shown to be required for calcineurin activation upon mycobacterial infection, thereby blocking their lysosomal delivery: when macrophages from TACO-deficient mice were studied, calcineurin was not activated, resulting in lysosomal transfer and death of internalized mycobacteria [7]. Although the prevailing paradigm states that *M. tuberculosis* prevents

lysosomal fusion, and persists and replicates within the phagosomes of macrophages, a recent study examining an extended time course for up to 7 days of infection arrived at a different conclusion: van der Wel et al. [9] report the escape of mycobacteria into the cytoplasm, which they consider a pathogenic feature of virulent mycobacteria. However, this is contradicted by similar ultrastructural studies on different mycobacterial species, which were found to remain within phagosomes even at late times of infection [10].

The interaction between *M. tuberculosis* and its host cells is very complex and, although it has been very extensively studied, is far from being understood. The role of the innate immune receptors involved in *M. tuberculosis* recognition and binding, in particular, has been mostly based on in vitro examination in transfected cells; studies using inhibitors or animals deficient in specific receptors have indicated that these receptors can compensate for each other or are dispensable [11–13]. This review will focus on the role of membrane-bound (as opposed to soluble) innate immune receptors of phagocytes (fig. 1) that have been shown to recognize mycobacteria (i.e. ligands expressed on the surface of mycobacteria), in a non-opsonic fashion. In contrast to non-opsonic binding, opsonic receptors recognize microbes that are coated with various proteins as target molecules for recognition such as antibodies, complement proteins, lectins and surfactants. It is generally accepted that non-opsonic binding is important during primary infection with inhaled bacteria as serum and complement components are limited in the alveolar space [14]. However, the role of surfactant-opsonized mycobacteria in phagocyte binding is not insignificant [15]. Opsonized or serum-mediated uptake

of *M. tuberculosis* may be more important at later stages of the infection [16, 17]. Also, cytoplasmic innate receptors such as nucleotide-binding oligomerization domain (NOD)-like receptors are important for innate immune recognition of *M. tuberculosis* [reviewed in ref. 18]. A recent study showed that macrophages and dendritic cells derived from NOD2-deficient mice are impaired in the production of pro-inflammatory cytokines upon mycobacterial infection compared to wild-type cells. However, no differences were observed in vivo with regard to susceptibility to *M. tuberculosis* infection [18].

In the present article, we will particularly concentrate on phagocytic membrane receptors involved in the very first line of defence against inhaled mycobacteria. These receptors are germ-line encoded and are also referred to as 'pattern recognition receptors' (PRRs), which recognize highly conserved molecular patterns (pathogen-associated molecular patterns) found only in microorganisms. The role of these receptors in *M. tuberculosis* infection, particularly their interaction with mycobacteria, downstream effector mechanisms, and their relevance in vivo will be discussed.

C-Type Lectins

Carbohydrate-binding C-type lectins play an important role in mycobacterial binding and in driving inflammatory responses due to the presence of carbohydrate-rich mycobacterial surface molecules. C-type lectins comprise a large family of proteins, divided into 17 groups, which contain one or more structurally related C-type lectin-like domains [reviewed in ref. 19]. One distinguishes between soluble lectins and cell-associated (or transmembrane) C-type lectins. Of the soluble C-type lectins, pulmonary surfactant proteins A and D, the collectins SP-A and SP-D, are important for mycobacterial infection: while SP-A increases *M. tuberculosis*-macrophage interaction by both directly upregulating phagocytosis and serving as a bacterial opsonin, SP-D agglutinates *M. tuberculosis* and decreases phagocytosis [reviewed in ref. 15]. Also, the soluble mannose-binding lectin which is primarily found in blood, was found to enhance infection of *M. tuberculosis* by facilitating the entry of mycobacteria into phagocytes, pathogen spread, and the establishment of infection [15]. SP-A, SP-D and mannose-binding lectin generally serve as opsonins and are therefore not the focus of this review.

Of the transmembrane C-type lectins, the MR, DC-SIGN and dectin-1 are most important for infection with

M. tuberculosis and will be discussed below. Although complement receptor-3 (CR3), another important macrophage receptor for mycobacterial infections, theoretically belongs to the integrin superfamily, it contains a lectin site that interacts with mycobacterial ligands and will be discussed in this section.

The Mannose Receptor

The MR (CD207) is a type I transmembrane glycoprotein with a large extracellular region containing 8 linear carbohydrate recognition domains (CRDs), and a short cytoplasmic tail containing a non-canonical tyrosine-based motif involved in phagocytosis, endocytosis and endosomal sorting [20]. The CRDs recognize a broad spectrum of ligands, including mannose-, N-acetylglucosamine- and fucose-terminated glycoconjugates [21, 22]. The MR is expressed predominantly on alveolar macrophages as well as monocyte-derived macrophages and dendritic cells, and represents a prototypic PRR recognizing the most abundant mycobacterial lipoglycan, mannose-capped lipoarabinomannan (ManLAM) [23]. Indeed, phagocytosis of *M. tuberculosis* by human macrophages occurs primarily via the MR [24]. However, this association depends on the length and abundance of surface-exposed ManLAMs, and the binding and phagocytosis of mycobacteria by the MR is limited to certain virulent strains of the *M. tuberculosis* complex [24]. It has been shown that higher-order phosphatidyl-myo-inositol mannosides (PIMs), particularly tri-acylated forms, as well as arabinomannans, mannans and mannoproteins can bind to this receptor [25].

It is still not clear whether the MR possesses a signalling function as it lacks defined signalling motifs in its cytoplasmic tail [20]. The binding of *M. tuberculosis* to the MR via ManLAM does not result in killing of the bacteria, but rather triggers anti-inflammatory pathways by interfering with the lipopolysaccharide-induced positive signals delivered by the TLRs such as IL-12 production [26]. Moreover, engagement of the MR by ManLAM during phagocytosis was shown to be crucial in limiting phagosome-lysosome fusion, allowing the bacteria to reach their unique niche within the host cell, the mycobacterial phagosome [27]. Anti-inflammatory and suppressive mechanisms are also triggered upon mycobacterial engagement of DC-SIGN via ManLAM. The relative importance of MR versus DC-SIGN ligation is briefly discussed below. The biological role of the MR in mycobacterial infections still has to be investigated in *M. tuberculosis*-infected MR-deficient mice.

Dendritic Cell-Specific Intercellular Adhesion Molecule-3 Grabbing Nonintegrin

DC-SIGN (CD209) is a type II mannose-binding C-type lectin receptor with a single extracellular CRD that clusters to form tetramers which are required for high-affinity binding of ligands [28]. Its intracellular region contains three motifs: a tyrosine-based internalization motif as well as a tri-acidic amino acid cluster and a di-leucine motif which are thought to be involved in phagocytosis and intracellular trafficking of ligand particles which can ultimately lead to phagosome-lysosome fusion [15].

DC-SIGN is expressed primarily on dendritic cells but has also been found on discrete subsets of macrophages [29] and has been shown to be induced on alveolar macrophages upon mycobacterial infection [30]. Initially described as a receptor for HIV gp120, DC-SIGN has been shown to recognize various microbial pathogens including viruses, bacteria, fungi and parasites [31]. DC-SIGN is not to be considered a classical PRR, but rather serves as an adhesion molecule which can be co-opted by microorganisms to their own advantage, leading to inhibition of the immunostimulatory function of dendritic cells and, hence, to the promotion of pathogen survival [32].

Recent studies revealed the importance of DC-SIGN as the major phagocytic receptor for *M. tuberculosis* on human dendritic cells by binding to ManLAM [33, 34]. Although some typical mycobacterial macrophage receptors are also expressed on dendritic cells such as the MR or CR3, they seem to be neglected by the tubercle bacillus [34].

It has been demonstrated that DC-SIGN discriminates between *Mycobacterium* species through selective recognition of the mannose caps on LAM [35]. DC-SIGN binds particularly strongly to *M. tuberculosis* and *M. bovis* BCG, while mycobacterial strains lacking ManLAM such as the avirulent *M. smegmatis* are not bound by this receptor [33]. However, DC-SIGN can also bind mycobacterial lipomannan (LM), arabinomannan, the 19-kDa antigen, and lower- and higher-order PIMs [25, 33].

Binding of mycobacteria by DC-SIGN results in an anti-inflammatory immune response by blocking maturation of infected dendritic cells through secreted ManLAM and inducing IL-10 production [33], enabling the intracellular survival of these organisms but not their replication [36]. It is believed that DC-SIGN cooperates with TLR4, and that binding of ManLAM by DC-SIGN triggers signals that interfere with the mycobacterial-induced dendritic cell maturation signals presumably generated by TLR4 [33]. Suppression of the protective pro-inflammatory immune response by *M. tuberculosis* binding to DC-

SIGN was recently confirmed in a study addressing genetic polymorphisms in the human *CD209* promoter region: a single-nucleotide polymorphism (SNP) at position -336A/G led to downregulation of *CD209* mRNA expression which was associated with significant protection against tuberculosis in the studied individuals [37]. However, this contradicts an earlier study that reported that the two promoter variants -871G and -336A decrease the risk of developing tuberculosis [38]. This discrepancy might be due to the different populations studied.

Although macrophages rather than dendritic cells are the primary target for mycobacterial infection, the specific function of dendritic cells in the cellular immune response seems to be modulated by *M. tuberculosis* [26]. Moreover, the intracellular behaviour of *M. tuberculosis* within dendritic cells differs from that in macrophages, with failure to replicate, which is linked to its different portal of entry [36]. It is possible that mycobacterial phagosomes mature to late endosomal/lysosomal stages in dendritic cells as opposed to macrophages where mycobacteria arrest phagosome maturation at an early endosomal stage, which promotes mycobacterial growth [32]. Indeed, it has been demonstrated that DC-SIGN directs phagocytic particles to the phagolysosome in transfected COS-1 fibroblast cells [27]. The observation that macrophages serve as the major intracellular niche to *M. tuberculosis* as opposed to dendritic cells could be explained by the different trafficking fate of *M. tuberculosis* after phagocytosis in the two cell types, which is linked to the receptor used: macrophages express high levels of MR while dendritic cells express high DC-SIGN activity, the latter being more potent in processing endocytosed products in lysosomes [27]. Thus, it is more favourable for the survival of the tubercle bacilli to enter the host via macrophages.

M. tuberculosis infection studies of mice lacking the murine DC-SIGN homologue SIGNR1 revealed no difference between wild-type and knockout animals in terms of survival rates and mycobacterial loads in the lungs or distant organs [39]. However, cellular expression and function of DC-SIGN and SIGNR1 are significantly different as SIGNR1 is not present on alveolar macrophages in uninfected mice nor is it induced on lung macrophages during infection. Although SIGNR1 and DC-SIGN display similar binding specificity, the role of SIGNR1 seems to be limited in murine *M. tuberculosis* infection [39].

Dectin-1

Dectin-1 is a type II transmembrane receptor which contains a single extracellular CRD on a stalk, and a cy-

toplasmic tyrosine-based activation-like motif which is involved in cellular activation [reviewed in ref. 40]. It is expressed on myeloid cells such as macrophages, dendritic cells and neutrophils, and a subset of T cells. Dectin-1 acts as a PRR and recognizes fungal-cell-wall-derived β -1,3-glucans in a calcium-independent manner and mediates numerous cellular responses, including microbial uptake and killing as well as the production of several cytokines and chemokines, and can directly couple innate and adaptive immune responses [41].

Extensively investigated as a receptor for fungal pathogens, the role of dectin-1 and its associated signalling pathways in any bacterial infection has not been demonstrated until recently when several studies started to point towards a participation of dectin-1, probably in cooperation with TLR2, in the pro-inflammatory response to infection with mycobacteria [42–44]. Murine bone-marrow-derived macrophages infected with avirulent or attenuated mycobacteria such as *M. avium*, *M. smegmatis*, *M. phlei* or *M. bovis* BCG showed a dectin-1-dependent production of TNF- α , IL-6, and G-CSF as opposed to virulent *M. tuberculosis* H37Rv [42]. The authors speculate that the limited stimulation of dectin-1 by virulent mycobacteria might be partly responsible for the minimal macrophage pro-inflammatory response induced by these mycobacteria in contrast to avirulent mycobacteria [45]. However, this is controversial as some investigators find no deficiency in inflammatory responses induced by virulent mycobacteria although these responses vary from strain to strain [46, 47].

A study on splenic dendritic cells showed that dectin-1 triggers the production of IL12p40 and IL12p70 independently of TLR2 upon infection with *M. tuberculosis* through an unknown mechanism involving Syk signalling [43]. Using the non-tuberculous mycobacterium *M. abscessus*, it could be demonstrated that dectin-1 in direct cooperation with TLR2, was required for phagocytosis of this organism in murine macrophages with subsequent activation of Syk and production of TNF- α , IL-6 and IL12p40 [44]. Although the interaction of dectin-1 with TLR2 was not observed in all of these studies, the collaboration of both receptors and the subsequent pro-inflammatory cytokine production by macrophages does occur via the Syk kinase pathway [48].

While the participation of dectin-1 in mycobacterial infection has begun to be elucidated, it is still not clear what ligand(s) on mycobacteria interact with dectin-1. The presence of β -glucans has not been described in mycobacteria but it is known that *M. bovis* BCG and *M. tuberculosis* express α -glucan within the outer capsule [49].

Also, dectin-1 might not be restricted to the recognition of β -glucans as it can also bind an endogenous but undefined ligand on T cells [50]. It has been shown that live mycobacteria bind directly to dectin-1 in a laminarin-inhibitable manner [42–44], indicating the presence of a ligand for dectin-1 on the bacterial envelope, which might mimic carbohydrate structures present in fungal β -glucans. Inhibition of mycobacterial binding and/or cytokine production by laminarin has also been shown in studies on CR3 (see paragraph below).

Taken together, the function of dectin-1 in host defence upon infection with mycobacteria is still unclear and probably dependent on the cell type being infected and the mycobacterial strain used. In vivo studies on dectin-1-deficient mice are urgently needed to define the biological role and regulatory mechanisms of this receptor for mycobacterial infection.

Complement Receptor 3

M. tuberculosis has been shown to activate the alternative complement pathway and become opsonized by C3b and iC3b which then enables interaction with complement receptors CR1 and CR3/CR4 [51]. Moreover, a C2a-dependent entry pathway specific for mycobacteria has been described [52]. It should be mentioned here that CR3 is the major complement receptor involved in phagocytosis of mycobacteria as this receptor mediates approximately 80% of complement-opsonized *M. tuberculosis* phagocytosis [51]. Interestingly, CR3 is able to mediate both complement-opsonized as well as non-opsonized phagocytosis of micro-organisms by different mechanisms, both of which have been described for mycobacteria [reviewed in ref. 53].

CR3, also named integrin $\alpha_M\beta_2$ or CD11b/CD18 or Mac-1, is a heterodimeric surface receptor that belongs to the integrin superfamily. It is expressed on neutrophils, monocytes, natural killer cells and alveolar macrophages. However, resident alveolar macrophages have been shown to express CR3 poorly while differentiation of alveolar macrophages in vitro resulted in increased expression of CR3, which was accompanied by an increased capacity to bind mycobacteria [17]. CR3 displays a broad ligand spectrum due to multiple binding sites; those ligands include complement fragment iC3b, intracellular adhesion molecule-1 (ICAM-1), and bacterial products [reviewed in ref. 53].

It has been shown that mycobacterial antigen 85C is recognized by the I domain of CR3, which also binds to complement molecule iC3b, while the lectin-like C-domain of CR3 binds mycobacterial oligosaccharides such

as LAM, which probably is the predominant binding site for mycobacteria [53]. This lectin-like domain has been characterized as a binding site for β -glucans as well as mannose, glucose and N-acetyl-D-glucosamine [54]. Indeed, competition studies showed that the binding of *M. tuberculosis* was inhibited by the β -glucan laminarin although this might simply represent the broad ligand spectrum of CR3 [53]. Studies on dendritic cells isolated from CD11b $^{-/-}$ mice showed that laminarin-blocked *M. tuberculosis* triggered IL12p40 production independently of this subunit of CR3 [43].

Binding of mycobacterial PIMs to CR3 has been described as well [55], suggesting that the receptor's lectin site might recognize the mannosyl moiety of PIM. The binding of mycobacteria to the lectin-like domain of CR3 could be blocked by β -glucans, which resulted in a strong inhibition of phagocytosis [53]. Differences in oligosaccharide composition influence the capability of various mycobacterial strains to bind to the lectin-like domain of CR3: studies on CR3-transfected cells revealed that pathogenic mycobacteria, but not non-pathogenic species, were internalized by this receptor [53]. However, phagocytosis of mycobacteria by CR3 does not induce killing of the bacteria [53].

Studies on macrophages isolated from CR3-deficient mice revealed that this receptor plays a role in the uptake of viable *M. tuberculosis*. However, CR3 does not play any role in the induction of microbicidal mechanisms and subsequent survival of the bacteria [56]. The observation that CR3-deficient and wild-type mice are equally resistant to *M. tuberculosis* infection suggests that in the absence of CR3, phagocyte infection by mycobacteria can be efficiently supported by alternative receptors [12]. Therefore, the biological role of CR3 in mycobacterial infection in vivo remains uncertain.

Toll-Like Receptors

The TLRs belong to an evolutionary highly conserved family of membrane receptors that have homology to the *Drosophila* Toll system. The mammalian TLR family now comprises 11 members which recognize a variety of microbial products including bacterial LPS, lipoteichoic acid, peptidoglycans, unmethylated CpG motifs of bacterial DNA or double-stranded RNA of viruses. They are expressed on various immune and non-immune cells including macrophages and dendritic cells [reviewed in ref. 57]. It has recently been demonstrated that the expression of various TLRs is upregulated in whole-blood samples of

patients with active pulmonary tuberculosis [58]. The TLRs are type I transmembrane proteins that contain repeated leucine-rich motifs in their extracellular domain which are involved in microbial recognition while their cytoplasmic domains are homologous to the signalling domain of the IL1 receptor. The interaction of *M. tuberculosis* with TLRs leads to phagocyte activation and may modulate pathways of phagocytosis, but does not directly initiate ingestion of the microbes.

Indeed, activated TLRs trigger intracellular signalling cascades [reviewed in ref. 59], most of which relying heavily on the adapter protein MyD88 and leading to pro-inflammatory and antimicrobial innate immune responses such as NF- κ B activation, MAP kinase activation and production of TNF- α , IL-1, IL-12, chemokines and nitric oxide. It is generally believed that a dominance of TLR signalling favours inflammatory and protective pathways as opposed to LAM signalling via C-type lectins (such as MR and DC-SIGN), which favours anti-inflammatory and suppressive mechanisms (discussed in [32]). Moreover, a recent study showed that the LM acylation pattern determines the pro-inflammatory versus anti-inflammatory effects of LM through different PRRs, i.e. TLR2-dependent versus TLR2-independent regulation [60]. However, some of the TLR-induced signals are anti-inflammatory and may mediate feedback inhibition to limit macrophage activation and prevent excessive inflammation: a recent study demonstrated that mycobacterial early secreted antigenic target protein-6 dampened TLR signalling by preventing assembly of the cytosolic MyD88-dependent signalling scaffold [61]. Further, TIR8, a member of the IL-1 receptor family, has recently been shown to play a key role in dampening inflammation and tissue damage in *M. tuberculosis* infection: TIR8-deficient mice succumbed rapidly to infection with *M. tuberculosis* [62].

As TLRs themselves do not function as phagocytic receptors, they are often complemented or even rely on other PRRs such as C-type lectins and scavenger receptors for efficient activation and ligand binding [reviewed in ref. 63]. Cross-talk between various receptors including TLR2, TLR4 and the MR for efficient IFN- γ production upon mycobacterial binding has recently been described in a tuberculous pleurisy model [64].

Mycobacteria can activate TLR2, TLR4, TLR9 and TLR1/TLR6 that heterodimerize with TLR2 and have all been implicated in recognition of mycobacterial antigens [reviewed in ref. 65]. Mycobacterial induction of TNF- α and IL-12 production was found to be primarily dependent on TLR2 signaling and to a lesser extent TLR4 signalling [66], TLR2 and TLR9 being known to regulate

dendritic-cell-derived IL-12 production in murine *M. tuberculosis* infection [67]. It has been shown that TLR2 recognizes mycobacterial lipoproteins such as the 19-kDa antigen (LpqH), LprA (Rv1270) and LprG (Rv1411) as well as LM and PIMs, while TLR2-dependent cell activation by LAM via phosphoinositide-capped lipoarabinomannan and arabinose-capped lipoarabinomannan has only been described for fast-growing mycobacteria [reviewed in ref. 59]. Cells are activated through TLR4 by binding to a heat-labile factor associated with *M. tuberculosis* as well as LM [5] and PIM [68], while TLR9 can sense the presence of *M. tuberculosis* DNA [67]. Man-LAM which is mainly recognized by the MR and DC-SIGN, resulting in anti-inflammatory effects, is not recognized by any TLR, suggesting an interesting correlation between LAM-capping structure and its immunomodulatory effects [59].

Although studies on mice infected with *M. tuberculosis* aerosol demonstrated a role for TLR2 and TLR4 in long-term control of the infection [68, 69], TLR2 and TLR4 seemed to be redundant in controlling *M. tuberculosis* infection in another mouse model: only at high infectious doses was reduced survival reported in TLR2-deficient mice [70]. Similar results were obtained in a study using *M. avium*-infected TLR2- or TLR4-deficient mice [71]. Infected mice deficient in both TLR2 and TLR9 revealed a major cooperation between these two receptors for host resistance to mycobacteria [67]. However, a recent study on mice infected with *M. tuberculosis* aerosol deficient in all TLRs to which *M. tuberculosis* products have been assigned as ligands, i.e. TLR2, TLR4 and TLR9, demonstrated that expression of pro-inflammatory cytokines was similar to that in wild-type animals. Also, both wild-type and TLR2/4/9 triple-deficient mice were able to control *M. tuberculosis* replication, suggesting that TLR2, TLR4 and TLR9 are not critical for triggering macrophage effector mechanisms central to anti-mycobacterial defence [72]. TLR2 forms heterodimers with either TLR1 or TLR6, and studies on TLR6-deficient mice showed that these animals are also resistant to high-dose *M. tuberculosis* aerosol infection [73]. The role of TLR1 in in vivo host response has not been determined yet. Interestingly, studies using mice deficient in the adapter molecule MyD88, which is common to signalling by most TLRs, but also IL1 receptor, succumbed to unrestrained mycobacterial growth while it was not required for the induction of adaptive T cell responses [72].

Taken together, the role of TLRs in immune responses to *M. tuberculosis* in vivo remains controversial and strongly depends on the experimental setting such as differences

in the mycobacterial strains used, differences in the nature and the dose of the microbe, and differences in genetic backgrounds of the receptor-deficient mice. However, TLRs seem to be relevant in human immunity to tuberculosis as a SNP in the human *TLR2* gene was associated with higher susceptibility to tuberculous meningitis [74].

Scavenger Receptors

Scavenger receptors belong to a broad family of cell surface transmembrane glycoproteins with multidomain structures classified into 6 subgroups based on their tertiary structure. Initially described to be important for atherogenesis they also play important roles in innate immunity and macrophage regulation. SRs display broad ligand-binding abilities, including recognition of modified lipoproteins, polyanionic molecules as well as Gram-positive and Gram-negative bacteria [reviewed in ref. 75].

Although the role of SRs in non-opsonic recognition and binding of mycobacteria has not been extensively studied, there are a few reports that point towards an implication of SRs in these mechanisms. Initially, class A SRs have been suggested to be important receptors for *M. tuberculosis* on human monocyte-derived macrophages: when binding to CRs and the MR was blocked, all binding that still persisted could be abrogated when class A SRs were competitively blocked with fucoidin and dextran sulfate [13]. Adipocytes which express numerous SRs can also display macrophage-like characteristics and phagocytose particles. Binding of *M. tuberculosis* to adipocytes of murine or human origin could be blocked by SR ligands such as polyinosinic acid, fucoidan and oxidized low-density lipoprotein species [76]. Recently, a genome wide RNAi screen for *Drosophila* macrophage-like cells identified Peste (Pes), a member of the CD36 family of SRs, to be essential for entry of *M. fortuitum* [77]. Moreover, human class B scavenger receptors were shown to mediate uptake of *M. fortuitum*, while murine CD36 did not [77].

Although no signal-transducing activity has been attributed to any type of SR, it has recently been shown that CD36 can augment TLR2 signalling in response to a subset of TLR2 ligands [78].

CD14

CD14 is a cell surface lipid-anchored glycan-linked protein without transmembrane and cytoplasmic domains which is predominantly expressed on myelomonocytic cells.

cytic cells. It mainly binds LPS of Gram-negative bacteria but can also interact with lipoteichoic acids and peptidoglycan of Gram-positive bacteria [79]. Further, CD14 can bind mycobacterial LAM, which leads to the secretion of IL-8 by macrophages [80]. It has also been shown that *M. tuberculosis* chaperonin 60.1 partially activates cells via a CD14-dependent mechanism [81]. Earlier studies demonstrated that CD14 can mediate uptake of *M. tuberculosis* by human microglial cells [82]. Generally, binding of microbial products by CD14 ultimately leads to phagocytosis, cellular activation and cytokine secretion, but requires cooperation with other receptors and transducer elements such as TLRs [83]. It has been shown that the concentration of soluble CD14 (which is produced by enzymatically cleaved membrane CD14) is significantly increased in patients with pulmonary tuberculosis [84]. Moreover, an SNP in the promoter region of the human CD14 gene, which is associated with elevated levels of soluble CD14 was associated with a higher risk of developing pulmonary tuberculosis [85].

Studies on CD14-deficient murine bone-marrow-derived macrophages demonstrated a significant decrease in TNF- α production upon infection with *M. avium* compared to control wild-type mice. In vivo infection of these CD14-deficient mice with *M. tuberculosis* or *M. avium* revealed no difference in mounting a protective and granulomatous response to mycobacterial infection compared to wild-type control mice [22, 70]. However, a recent study described a role for CD14 in chronic *M. tuberculosis* infection. Whilst no difference in bacterial load in the lung was found between wild-type and CD14-deficient mice up to 32 weeks after high-dose infection, the wild-type mice started to succumb to infection from 20 weeks onwards while the CD14-knockout mice did not [86]. Therefore, CD14 deficiency seems to protect the animals from chronic *M. tuberculosis* infection via a reduction in the inflammatory response.

Other Receptors

Other receptors such as CD40, CD43 and CD44, which have been shown to play a role in mycobacterial recognition will be briefly summarized here.

CD40, a member of the TNF receptor family and a costimulatory molecule on antigen-presenting cells, has been shown to be involved in *M. tuberculosis* recognition via hsp70 expressed on mycobacteria [87] leading to stimulation of CC-chemokine production. Another study suggested that CD40 is not directly involved but drives

inflammation and enhances immune responses to *M. tuberculosis* infection in vivo when dendritic cells are stimulated via CD40 although the mycobacterial burden in the lungs is not reduced compared to control animals [88].

CD43, a transmembrane sialoglycoprotein expressed on most hemopoietic cells, was initially studied in vitro in CD43-deficient macrophages, revealing a role of this surface mucin in binding and/or uptake of mycobacteria and in mycobacterium-induced TNF- α production [89]. This CD43-mediated induction of pro-inflammatory cytokines is thought to regulate apoptosis in order to control intracellular growth of mycobacteria [90]; aerosol-infected CD43-knockout mice showed an increased mycobacterial load during both the acute and chronic stage of infection and a more rapid development of granulomas compared to wild-type animals.

CD44, an adhesion molecule which is upregulated during inflammatory response, has also been shown to be a receptor for *M. tuberculosis* on macrophages that mediates phagocytosis [91]. However, CD44-deficient mice infected with *M. tuberculosis* showed neither a difference in pulmonary bacterial load compared to wild-type animals nor an altered expression of Th1 immunity to tuberculosis [92]. Nevertheless, CD44 plays an important role in controlling and regulating lung inflammation as absence of this molecule resulted in a substantially increased accumulation of neutrophils in the lung [92].

Cholesterol

Although cholesterol is not a receptor, its role in mycobacterial uptake has emerged in recent years and it is suggested that cholesterol might be directly involved in receptor-mediated phagocytosis. Therefore, studies on interactions between mycobacteria and cholesterol are mentioned below.

Mycobacteria have been demonstrated to display a high binding capacity for cholesterol compared to other microorganisms [2], possibly by direct binding to cholesterol of components of the extremely glycolipid-rich mycobacterial cell wall. Indeed, cholesterol accumulates at the site of *M. tuberculosis* entry into macrophages, and depleting cells of cholesterol prevents *M. tuberculosis* internalization [2], indicating that cholesterol accumulation around phagocytic receptors rather than the nature of the receptors itself may dictate *M. tuberculosis* uptake. Moreover, TACO, a component of the phagosome coat that prevents degradation of mycobacteria in lysosomes, was

shown to be associated with cholesterol-rich membrane regions [2]. Therefore, entering the cholesterol-rich domains of macrophages where TACO is predominantly found might ensure mycobacterial intracellular survival in TACO-coated phagosomes [2]. Moreover, cholesterol is also required for mycobacteria to prevent fusion of phagosomes with lysosomes: when pre-existing phagosomes infected with *M. avium* were depleted of cholesterol, they were reported to mature and fuse with lysosomes [93].

One of the receptors that might confer cholesterol-dependent non-opsonic mycobacterial uptake has been identified in neutrophils as being CR3. It was shown that upon infection with *M. kansasii*, CR3 and associated GPI-anchored proteins relocate in cholesterol-rich domains where internalization of the bacterium occurs [94].

It has been speculated that hypocholesterolaemia is a major risk factor for developing pulmonary tuberculosis [95], and patients receiving a cholesterol-rich diet showed an accelerated bacteriologic sterilization of sputum cultures. However, it has not yet been determined whether in vivo a similar relocation of cholesterol occurs at sites of *M. tuberculosis* entry as described in [2].

Another interesting association between cholesterol and infection with *M. tuberculosis* has been reported only recently: mice deficient in apolipoprotein E (ApoE) succumbed to a low-dose *M. tuberculosis* aerosol infection only 4 weeks after infection when fed a high-cholesterol diet compared to ApoE-deficient mice on a low-cholesterol diet or wild-type mice on either diet [96]. ApoE-deficient mice display a hypercholesterolaemic phenotype as they are unable to take up blood cholesterol. These highly elevated serum cholesterol levels when fed a high-cholesterol diet were found to be responsible for the extreme *M. tuberculosis* susceptibility of these mice. The underlying mechanism is not clear but *M. tuberculosis* can take up and use cholesterol as a source of energy and might therefore have a growth advantage due to the increased nutrient availability. This hypothesis was confirmed by a recent study showing that a mycobacterial gene cluster encoding genes involved in cholesterol catabolism is specifically expressed during survival of *M. tuberculosis* in macrophages [97].

Conclusion

Taken together, the interaction of mycobacteria with both opsonic and non-opsonic receptors on phagocytic cells is complex and incompletely understood. The role of

a particular receptor is often studied in vitro in an artificial system and isolated from the physiological context while data obtained from in vivo studies are often very controversial and the results seem to vary largely in different experimental settings. It is more than likely that in vivo, *M. tuberculosis* is not internalized by macrophages using a single receptor-mediated pathway. Mycobacteria display numerous and diverse ligands on their surface which might either engage multiple receptors of multiple types simultaneously, or might bind to those receptors that are available on a particular cell or cell type. Moreover, mycobacterial surface molecules might also function to impair the formation of adequate receptor clustering, i.e. cooperation of various receptors to cross-talk and coordinate recognition and entrance of mycobacteria into phagocytes. The role of cholesterol in mycobacterial uptake and survival has been discovered recently and further adds to the complexity of macrophage-mycobacterial interactions (fig. 1).

Various studies propose that progression of tuberculosis following infection is determined by the type of encounter between mycobacteria and host cell receptors such as C-type lectins versus TLR, and consequently the type of host cell such as macrophage versus dendritic cells. However, the question still remains whether *M. tuberculosis* or the receptor engaged by *M. tuberculosis* (or both) determine the fate of the phagosome and, consequently, the course of disease. As mentioned above, a dominance of TLR signalling would favour inflammatory and protective pathways while a dominance of signaling via C-type lectins would favour anti-inflammatory and suppressive/non-protective mechanisms. In this context, the LAM capping structure displays an interesting correlation to their immunomodulatory effect: Phosphoinositide-capped lipoarabinomannans are usually considered pro-inflammatory molecules stimulating the production of TNF and IL-12 while ManLAM are rather anti-inflammatory molecules stimulating IL-10 production and inhibiting IL-12 and TNF production [59].

It is possible that the different PRRs on alveolar macrophages outcompete LAM-mediated inhibitory signaling, thus favoring protective effector functions [32]. The overall effect probably depends on various parameters, especially the expression and availability of particular receptors and their triggered downstream responses.

As mentioned before, inconsistencies in the various studies on mycobacterium-receptor function might be due to several technical aspects. The substantial complexity in host defenses explains why specific components have been found to be either essential or redun-

dant. This is dependent on both the nature of the microbe and perhaps also the site of infection (i.e. cell type infected). For example, the phenotype of the macrophages (resident vs. elicited vs. immune activated) as well as the way how mycobacteria are prepared in a particular experimental setting (sonication vs. non-sonication) greatly influences phagocyte-mycobacterium interaction [16, 98]. It has been shown previously that it is important to use experimental infection procedures that most closely mimic natural exposure to obtain results that are physiologically most relevant: while dynamic (shear) binding conditions select more for higher-affinity interactions and imitate the physiological situation better, static binding of mycobacteria to cells would generate more non-specific background binding as demonstrated in a recent in vitro study [99]. In terms of the in vivo system, *M. tuberculosis* has been shown to be more

virulent when administered aerogenically rather than intravenously [100].

As adhesion and internalization are the first steps in mycobacterial pathogenesis, unraveling the role of the receptor(s) involved in these processes is of utmost importance. Inhibition of these events could constitute an efficient form of prophylaxis aimed at blocking invasion of *M. tuberculosis*.

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