**Mycobacterium abscessus** activates the NLRP3 inflammasome via Dectin-1–Syk and p62/SQSTM1

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Numerous atypical mycobacteria, including *Mycobacterium abscessus* (Mabc), cause nontuberculous mycobacterial infections, which present a serious public health threat. Inflammasome activation is involved in host defense and the pathogenesis of autoimmune diseases. However, inflammasome activation has not been widely characterized in human macrophages infected with atypical mycobacteria. Here, we demonstrate that Mabc robustly activates the nucleotide binding and oligomerization domain-like receptor family pyrin domain containing 3 (NLRP3) inflammasome via dectin-1/Syk-dependent signaling and the cytoplasmic scaffold protein p62/SQSTM1 (p62) in human macrophages. Both dectin-1 and Toll-like receptor 2 (TLR2) were required for Mabc-induced mRNA expression of pro-interleukin (IL)-1β, cathelicidin human cationic antimicrobial protein-18/LL-37 and β-defensin 4 (DEFB4). Dectin-1-dependent Syk signaling, but not that of MyD88, led to the activation of caspase-1 and secretion of IL-1β through the activation of an NLRP3/apoptosis-associated speck-like protein containing a caspase recruitment domain (ASC) inflammasome. Additionally, potassium efflux was required for Mabc-induced NLRP3/ASC inflammasome activation. Furthermore, Mabc-induced p62 expression was critically involved in NLRP3 inflammasome activation in human macrophages. Finally, NLRP3/ASC was critical for the inflammasome in antimicrobial responses to Mabc infection. Taken together, these data demonstrate the induction mechanism of the NLRP3/ASC inflammasome and its role in innate immunity to Mabc infection.

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The nontuberculous mycobacteria (NTM), also called atypical mycobacteria, are ubiquitous in soil and water and exhibit varied pathogenicity. NTM infections are now increasingly recognized as a cause of chronic lung disease, lymphadenitis, skin disease and disseminated infections.1,2 Rapidly growing mycobacteria such as *Mycobacterium abscessus* (Mabc), *M. fortuitum* and *M. chelonae* are the principal pathogens involved in NTM infections.1,2 Mabc (formerly *M. chelonae* subspecies *abscessus*) is one of the most common NTM species that causes invasive mycobacterial disease and disseminated infections in patients with cystic fibrosis.3,4 Antimicrobial peptides, including human cathelicidins and defensins, constitute a major component of innate host defense systems.5–7 Human cationic antimicrobial protein (hCAP-18)/LL-37 is the only member of the cathelicidin family to have been identified in humans.5,6 The small (3–5 kDa) human cationic defensins are a delineated family of effector molecules that contribute to host defense, inflammation and cytotoxicity.7 Mabc infection is often notoriously difficult to eradicate,8 however, the mechanisms of innate immune defense in host cells against Mabc infection remain unknown.

Nucleotide binding and oligomerization domain-like receptors (NLRs) are pattern-recognition receptors that recognize bacterial products in the cytosol.9 Recent studies have revealed that these receptors can activate caspase-1 via the inflammasome, a multiprotein complex.9,10 Upon cellular infection or stress, inflammasome activation has been shown to be important in the maturation of pro-inflammatory cytokines such as interleukin (IL)-1β, thereby triggering innate immune responses and inflammation.11–13 Accumulating studies have highlighted the roles of several pathways that activate the inflammasome.10,12 Recent studies have revealed that the cytoplasmic protein p62/SQSTM1 (p62) functions as a nodal point in various signaling pathways, including those that control inflammasome activation.13–15 In *Shigella*-infected cells, p62 is recruited to polyubiquitinated membrane remnants that are associated with inflammasome components and caspase-1.13 However, the possible role of p62 in the
activation of the inflammasome in innate immune cells has not been characterized. Additionally, the ability of atypical mycobacteria to activate the inflammasome and its regulation in monocytes/macrophages remain to be clarified.

Dectin-1, a C-type lectin-like pattern-recognition receptor present on monocytes/macrophages and dendritic cells, can recognize β-glucans. Dectin-1 contains an immunoreceptor tyrosine-based activation motif-like motif in its cytoplasmic tail, and when phosphorylated upon ligand binding, it can activate the spleen tyrosine kinase (Syk)-dependent pathway. 

Recent studies have demonstrated the role of dectin-1 in the activation of the inflammasome in fungal infections, including those with Candida albicans and Aspergillus fumigatus. Moreover, a recent study has shown that the dectin-1/Syk signaling pathway is involved in β-glucan-mediated nucleotide binding and oligomerization domain-like receptor family pyrin domain containing 3 (NLRP3 also called NALP3 and cryopyrin) inflammasome activation. However, the role of dectin-1 in inflammasome activation in mycobacterial infection, particularly in human cells, has not been fully characterized.

Our previous studies have shown that Mabc is actively internalized and induces proinflammatory cytokine secretion in macrophages through dectin-1-dependent signaling and cooperation between Toll-like receptor (TLR) 2 and dectin-1. In the present study, we investigated how Mabc activates the inflammasome, a process that is required for efficient innate immune activation in response to Mabc.

We show that Mabc activates the NLRP3/ASC inflammasome via dectin-1/Syk-dependent signaling. Both dectin-1 and TLR2 were required for the induction of pro-IL-1β, cathelicidin hCAP-18/LL-37 mRNA expression. Notably, dectin-1-dependent Ca2+ influx and Syk signaling contributed to the activation of the NLRP3/ASC inflammasome. Additionally, p62 expression was critical for Mabc-induced proinflammatory responses and inflammasome activation in human macrophages. Finally, NLRP3/ASC inflammasome activation contributed to antimicrobial responses against Mabc infection in human macrophages.

**RESULTS**

Mabc robustly induces caspase-1 activation and IL-1β maturation in human macrophages

We first examined whether Mabc activates caspase-1 and IL-1β secretion in human primary monocyte-derived macrophages (MDMs) at various time points after infection and at various moi (multiplicities of infection). When MDMs were treated with Mabc, IL-1β levels were increased significantly in culture supernatants beginning 3 h after infection (at moi of 1 and 3), and peaked at 18 h (Figure 1a). Because there was no significant Mabc-dependent IL-1β secretion between moi of 3 and 10, cells were infected at a moi of 3 in subsequent experiments (data not shown). Given that caspase-1 (an IL-1β-converting enzyme) cleaves the IL-1β precursor to mature IL-1β, we investigated caspase-1 activation and IL-1β processing in human macrophages using western blot analysis. After 3 h of Mabc infection, MDMs displayed clear activation of caspase-1, as evidenced by an increased amount of the cleaved p10 subunit (Figure 1b, bottom). Consistent with caspase-1 activation, the processed active form of IL-1β was evident in culture supernatants after 6 h of Mabc infection (Figure 1b, bottom). Additionally, IL-1β secretion after Mabc infection was considerably reduced when human MDMs were pretreated with a caspase-1-specific inhibitor, z-YVAD-fmk (Figure 1c). Together, these data suggest that Mabc robustly activates the inflammasome and induces IL-1β secretion in human macrophages.

**Dectin-1 and TLR2 have essential roles in the expression of pro-IL-1β, cathelicidin hCAP-18/IL-37 and DEFB4 in macrophages infected with Mabc**

Previously, we showed that dectin-1 is a receptor for Mabc recognition and cooperates with TLR2 to induce inflammatory cytokines in response to Mabc infection. In the current study, we examined the time course of the mRNA expression of LL-37 and DEFB4 in human macrophages after Mabc infection. Mabc infection rapidly led to expression of LL-37 and DEFB4 mRNA in human primary MDMs (Supplementary Figure S1a). Levels of LL-37 and DEFB4 mRNA peaked...
within 6h of Mabc infection and then decreased (Supplementary Figure S1a).

To investigate whether dectin-1 is involved in Mabc-induced pro-IL-1ß mRNA expression, we performed dectin-1 small-interfering RNA (siRNA) transfection to THP-1 cells before infection. As shown in Figure 2a, knockdown of dectin-1 significantly attenuated Mabc-induced mRNA expression of pro-IL-1ß, LL-37 and DEFB4 in human monocytic THP-1 cells. Additionally, Mabc-induced expression of pro-IL-1ß, LL-37 and DEFB4 mRNA was significantly inhibited by pretreatment with a neutralizing dectin-1 antibody (Ab; Supplementary Figure S1b). Expression of these mRNAs was not modulated by an isotype-matched control (IC) monoclonal Ab (mAb; Supplementary Figure S1b).

We further investigated the role of TLR2 in Mabc-induced expression of pro-IL-1ß, LL-37 and DEFB4 in human macrophages. As shown in Figure 2b, specific hTLR2 gene silencing in human THP-1 cells resulted in a profound inhibition of Mabc-induced pro-IL-1ß, LL-37 and DEFB4 mRNA expression, whereas neither hTLR4 gene silencing nor control vectors demonstrated this effect. These data strongly suggest that dectin-1 and TLR2 have essential roles in the Mabc-induced mRNA expression of pro-IL-1ß, LL-37 and DEFB4 in human macrophages.

Dectin-1-dependent Syk signaling, but not MyD88, leads to activation of caspase-1 and secretion of IL-1ß in macrophages infected with Mabc

Dectin-1 activates innate signaling pathways through the adaptor molecule Syk kinase. Additionally, dectin-1 induces Ca2+ flux in dendritic cells through phospholipase C-γ2 activation. We further examined the role of Syk and Ca2+ signaling in Mabc-induced inflammasome activation. Either silencing of Syk by specific siRNA transfection or pharmacological inhibition of Syk significantly inhibited IL-1ß maturation as well as caspase-1 cleavage in response to Mabc infection in THP-1 cells (Figure 3a) and MDMs (Supplementary Figure S2a), indicating that Syk activity modulates caspase-1 activation and IL-1ß secretion in macrophages infected with Mabc. As a control, Mabc-induced tumor necrosis factor (TNF-α) production was not inhibited in MDMs pretreated with picetannol (Supplementary Figure S2b).

We next examined whether Mabc-dependent dectin-1 activation could induce intracellular Ca2+ influx in human monocytes. As shown in Figure 3b, Mabc treatment of THP-1 cells led to a significant induction of intracellular Ca2+ influx, which was considerably dampened by preincubation of these cells with a blocking anti-dectin-1 Ab. These data imply that Mabc-dependent dectin-1 activation could trigger Ca2+ signaling in these cells. We additionally found that pretreatment with the intracellular Ca2+-specific chelator 1,2-bis (2-aminophenoxy)ethane-N,N,N,N-tetracetic acid tetra-(acetoxy-methyl) ester (BAPTA-AM) significantly blocked Mabc-induced caspase-1 activation and cleavage and secretion of IL-1ß in human MDMs (Figures 3c and d). However, Mabc-dependent TNF-α production was not substantially modulated in MDMs pretreated with BAPTA-AM (Figure 3d, bottom). Silencing of MyD88 by specific siRNA transfection did not affect formation of mature IL-1ß in THP-1 cells after Mabc infection (Figure 3e). These data imply that dectin-1-dependent Ca2+ and Syk signaling, but not TLR2-dependent MyD88 signaling, contribute to the production of mature IL-1ß in macrophages infected with Mabc.

NLRP3 and ASC are involved in the maturation and secretion of IL-1ß by human macrophages after Mabc treatment

Recent studies have shown that NLRP3/ASC is required for caspase-1 activation and mature IL-1ß secretion by M. tuberculosis protein early-secreted antigenic target (ESAT)-6 in human macrophages. Thus, we determined the effects of NLRP3 and ASC on Mabc-induced IL-1ß secretion by human primary MDMs. As shown in Figure 4a, knockdown of NLRP3 or ASC via lentiviral short hairpin RNA (shRNA)-mediated RNA interference resulted in a decrease of >90% in the mRNA levels of NLRP3 or ASC. Based on enzyme-linked immunosorbent assay (ELISA), analysis of IL-1ß and IL-8 secretion, knockdown of NLRP3 and ASC caused a significant reduction in IL-1ß, but not IL-8, production by human MDMs (Figure 4a). Additionally, lentiviral shRNA specific to the NLRP3 (shNLRP3) or ASC (shASC) genes significantly inhibited mature IL-1ß secretion in human primary MDMs after Mabc infection, compared with secretion in MDMs transduced with nonspecific control shRNA lentiviral particles (shNS; Figure 4b). These results suggest that Mabc-induced secretion and maturation of IL-1ß is dependent on NLRP3 and ASC expression in human macrophages.

Mabc-induced IL-1ß secretion does not require LL-37, but require potassium efflux

It has been shown previously that human cathelicidin-derived peptide LL-37 induces IL-1ß release in lipopolysaccharide (LPS)-primed monocytes through activation of the P2X7 receptor. We examined whether Mabc-induced LL-37 is required for the caspase-1 activation and IL-1ß maturation in MDMs. To examine this, MDMs transduced with shRNA specific for LL-37 (shLL-37) or shNS were treated with Mabc. The silencing of LL-37 using shLL-37 did not affect IL-1ß processing or caspase-1 activation in human macrophages (Figure 5a),
indicating that LL-37 is not responsible for Mabc-induced activation of IL-1β processing and release in macrophages.

Potassium efflux is implicated in the activation mechanism for the inflammasome, because NLRP3 inflammasome activation can be triggered by low intracellular potassium, but prevented by culturing cells in high potassium-containing media.29 To determine whether Mabc-induced IL-1β secretion was inhibited by high extracellular potassium levels, Mabc-infected MDMs were exposed to high potassium-containing media, and then IL-1β release was determined. As shown in Figure 5b, Mabc-stimulated MDMs in the presence of elevated potassium demonstrated significantly decreased IL-1β levels, showing that elevated extracellular potassium suppressed the ability of Mabc to induce secretion of IL-1β. However, Mabc-induced secretion of IL-8 was largely unaffected (Figure 5b). Furthermore, mature IL-1β secretion and caspase-1 cleavage in response to Mabc infection were significantly inhibited by treatment of MDMs with high concentrations of potassium (Figure 5c). These data suggest that potassium efflux, but not LL-37, is required for Mabc-induced NLRP3/ASC inflammasome activation in human MDMs.

Mabc-induced p62 expression is involved in NLRP3 inflammasome activation in human macrophages

The autophagy adapter p62 can recruit and oligomerize important signaling molecules in cytosolic cellular speckles to control cell survival, apoptosis and inflammation.13–15,30 We next examined whether p62 was induced in response to Mabc infection in human macrophages. Reverse transcription polymerase chain reaction (RT–PCR) and immunoblot analysis revealed that p62 transcripts and protein were constitutively expressed in unstimulated human macrophages. After Mabc treatment, p62 transcripts and protein were markedly increased after 3 h, peaked at 12–24 h and gradually decreased until 48 h (Figure 6a; Supplementary Figure S3a; human MDMs and THP-1 cells, respectively).

To investigate whether p62 had a role in IL-1β secretion in macrophages, human THP-1 cells transfected with siRNA specific for p62 (sip62) or siNS were treated with Mabc. After 18 h, silencing of p62 significantly reduced Mabc-induced synthesis of IL-1β (Supplementary Figure S3b) and the release of the caspase-1 p10 fragment and mature IL-1β (Supplementary Figure S3c) in human macrophages.

Figure 3 Syk, but not MyD88, has a role in Mabc-induced caspase-1 activation and IL-1β secretion in human MDMs. (a, e) THP-1 cells were transfected with control nonspecific siRNA (siNS) or specific siRNAs against hSyk (siSyk, a) or hMyD88 (siMyD88, e). (c, d) Human MDMs were pretreated with BAPTA (5, 10 or 25 μM) for 45 min. Cells were then infected with Mabc (moi=3) for 18 h. (a, c, e) Western blots of cell lysates (cell) and supernatants (SN) from THP-1 cells (a, e) and MDMs (c) were probed with anti-caspase-1 (Casp1) Ab and anti-IL-1β Ab. (b) (Top) THP-1 cells were infected with Mabc (moi=3). (Bottom) THP-1 cells were pretreated with or without anti-dectin-1 Ab (α-Dectin-1) or the same concentration of an IC Ab before infection with Mabc (moi=3). The cells were then loaded with Fluo-4/AM and imaged by LSM510 confocal microscope objective lens at 5 s intervals. Changes are shown as mean fluorescence intensity from 15 cells per microscopic field over time. (d) IL-1β and TNF-α ELISA analysis. Data are from a representative of at least three independent experiments (d; mean values ± s.d. of triplicate samples). *P<0.05; ***P<0.001 versus control cultures. M, Mabc; SC, solvent control (0.1% DMSO); U, uninfected.
NLRP3 and ASC have critical roles in antimicrobial responses to Mabc in human macrophages

Recent studies have shown that inflammasome activation contributes to the antimicrobial defense against zmp1-deleted M. tuberculosis infection. Therefore, we decided to address whether NLRP3 or ASC affects the antimicrobial response to Mabc infection in human MDMs. Human primary MDMs transduced with shNLRP3, shASC or control shRNA, as well as untransduced MDMs, were infected with Mabc. After 4 h, all extracellular bacteria were removed by washing three times with warm phosphate-buffered saline. The cells were then cultured for the times indicated and intracellular bacteria were harvested and assayed for viability using colony forming unit (c.f.u.) assays. On day 3, intracellular bacterial viability was significantly increased in human MDMs transduced with shNLRP3 or shASC, compared with viability in untransduced and shNS-transduced cells (Figure 7). These data suggest that NLRP3 and ASC have important roles in human macrophage defenses against Mabc.

DISCUSSION

Mabc is a common and rapidly growing mycobacterium that causes human diseases, including chronic pulmonary disease and skin and soft tissue infections. The management and treatment of Mabc infections is often difficult, and the specific host factors involved in the antimicrobial immune response to Mabc, in contrast to tuberculosis, remain largely unknown. Previous studies have shown that dectin-1, a natural killer cell-receptor-like C-type lectin, is required for the recognition of Mabc, M. ulcerans, M. tuberculosis and other mycobacteria. The engagement of dectin-1 by mycobacteria initiates intracellular signaling that activates nuclear factor-κB and mitogen-activated protein kinases, leading to the production of proinflammatory cytokines in macrophages and keratinocytes. Recent studies have defined the roles of dectin-1 and Syk in the activation of the NLRP3 inflammasome, which participates in host antifungal defense against the dissemination of mucosal infection and mortality in vivo. However, whether atypical Mabc activates the inflammasome and how it is regulated in human macrophages have not been determined. In the present report, we demonstrate that dectin-1/Syk signaling has a critical role in Mabc-induced inflammasome activation and IL-1β secretion in human macrophages.

We demonstrated that both dectin-1 and TLR2 act as inducers of pro-IL-1β mRNA. Our data partially correlate with previous studies showing that TLR2 and dectin-1 control pro-IL-1β gene transcription, whereas NLRP3 and caspase-1 regulate the processing and maturation of IL-1β in macrophages infected with C. albicans. Many fungi and mycobacteria are recognized by both TLR2 and dectin-1, whose interactions have an important role in the induction of inflammatory responses. Dectin-1-dependent signaling pathways can promote the synthesis of cytokines through an intracellular immunoreceptor tyrosine-based activation motif in a Syk-dependent manner. In human macrophages, we have defined an essential role for Syk in caspase-1 activation and IL-1β secretion in response to Mabc infection. Moreover, we also revealed the importance of intracellular Ca2+ influx, as well as the NLRP3 inflammasome (NLRP3 and ASC), in controlling Mabc-mediated IL-1β secretion. Our data correlate with a recent study, demonstrating that the dectin-1/Syk signaling pathway has a vital role in fungus- or β-glucan-mediated NLRP3 inflammasome activation. Either genetic deletion or pharmacological inhibition of Syk selectively abrogated NLRP3 inflammasome activation by Candida infection, but not by inflammasome activators such as Salmonella typhimurium or the bacterial toxin nigericin. Additionally, IL-1β processing and release were found to be dependent...
IL-1β NLRP3/ASC inflammasome activation through Ca2+-dependent dectin-1/Syk pathway activation is important for Mabc-induced intracellular mycobacteria. Moreover, knockdown of both receptor activation and contributes to antimicrobial activity against activities in human monocytes/macrophages. TLR2-mediated vitamin D-induced phagosome–lysosome fusion and antimicrobial studies have shown that hCAP-18/LL-37 has an essential role in infection. These observations suggest that DEFB4 and LL-37 may for optimal antimicrobial activity against intracellular mycobacterial infection requires both TLR2/1-mediated IL-1β and vitamin D receptor activation and contributes to antimicrobial activity against intracellular mycobacteria. Moreover, knockdown of both DEFB4 and cathelicidin ablates TLR2/1-mediated antimicrobial responses, suggesting that both DEFB4 and cathelicidin expression are required for optimal antimicrobial activity against intracellular mycobacterial infection. These observations suggest that DEFB4 and LL-37 may have important roles in innate immune responses against Mabc infection through TLR2/dectin-1-dependent signaling.

The multidomain scaffold protein p62 functions as a signaling adaptor molecule in cellular signaling pathways, particularly in the regulation of nuclear factor-kB through interaction with TNF receptor-associated factor 6. The biological functions of p62 have been studied widely in osteoclastogenesis, inflammation, differentiation and obesity. However, the exact mechanisms by which p62 mediates the activation of caspase-1 and IL-1β maturation have not been clarified. Our data suggest that Mabc-induced p62 expression is required for caspase-1 activation and the secretion of mature IL-1β in human macrophages. The polyubiquitin-binding protein p62/SQSTM1 is a common component of protein aggregates and yields protein bodies after polymerization. p62 is involved in linking protein aggregates to the autophagic machinery, autophagosomes and lysosomal structures, and thus reduces the toxicity of mutant protein aggregates in the cytosol. The current study does not examine whether Mabc infection induces autophagy in human monocytes. Indeed, numerous bacteria can utilize several virulence factors or strategies to evade the autophagy system. We have found that p62 levels were increased significantly in human MDMs after Mabc infection (see Figure 6a and Supplementary Figure S4a), indicating that Mabc may cause cytoplastic accumulation of p62 protein aggregates and thus modify or inhibit autophagosome formation in human macrophages. Defective autophagy resulted in the upregulation of p62, which is responsible for increased nuclear factor-kB regulation, and promoted pathophysiological conditions, including cellular proliferation, tumorigenesis and chronic inflammation. When autophagy is inhibited, p62-positive inclusion-protein aggregates may lead to inflammasome activation and IL-1β secretion. Defective autophagy results in the enhanced generation of mitochondrial reactive oxygen species (ROS), which in turn leads to the activation of the NLRP3 inflammasome and IL-1β secretion. Future studies will clarify whether Mabc induces the inhibition of macroautophagy, the accumulation of p62 protein and increased cellular stress including mitochondrial ROS, thus contributing to the activation of the NLRP3 inflammasome machinery and caspase-1 activation.

**Figure 5** Potassium efflux, but not LL-37, is required for Mabc-induced IL-1β secretion in human MDMs. (a) Human MDMs were transduced with shNS or shLL-37 for 3 days before infection with Mabc (moi=3) for 18 h and then subjected to western blot analysis. (b, c) Human MDMs were pretreated with high K+ buffer (10, 50 or 150 mM) for 45 min before Mabc infection. (b) Supernatants were harvested at 18 h after Mabc infection and subjected to ELISA for IL-1β and IL-8. (a, c) Western blots of cell lysates (cell) and supernatants (SN) were probed with anti-caspase-1 (Casp1) Ab and anti-IL-1β Ab. Data are from a representative of at least three independent experiments (b; mean values ± s.d. of triplicate samples). The densitometry values for cleaved caspase-1 p10 and 17 kDa IL-1β were normalized to β-actin levels (a, bottom). ***P<0.001 versus control cultures. M, Mabc; SC, solvent control (0.1% DMSO); U, uninfected.
Using c.f.u. assays, we demonstrated that NLRP3/ASC is required for the antimicrobial response to Mabc infection. Accumulating evidence indicates the pivotal role of inflammasome activation in host defense against several types of infection. Indeed, *M. tuberculosis* is known to prevent inflammasome activation and IL-1β secretion. The *M. tuberculosis* zmp1 gene, which encodes a putative Zn2+ metalloprotease, is required for activation of the inflammasome and enhanced antimicrobial responses in infected lungs of mice and in macrophages. However, recent studies using wild-type and ASC-deficient mice have demonstrated that ESAT-6 system-1 (Esx1)-dependent inflammasome activation exacerbates mycobacterial infection without restricting bacterial growth, suggesting a detrimental effect of Esx1-mediated inflammation. ESAT, a 6-kDa mycobacterial virulence factor, promotes IL-1β secretion and caspase-1 activation through the perturbation of host cell membranes and the formation of an infection-inducible inflammasome complex. Additionally, neither NLRP3 nor caspase-1 significantly affected the restriction of *M. tuberculosis* in vivo using an experimental tuberculosis infection model.
Bacterial culture

Mabc-type strain ATCC 19977 was cultured as previously described.23 Bacteria were grown at 37 °C on Middlebrook 7H10 agar medium (Difco Laboratories, Detroit, MI, USA) supplemented with 10% OADC (oleic acid, albumin, dextrose, catalase) (BD Pharmingen, San Diego, CA, USA) and 0.05% Tween 80 (Sigma-Aldrich). Mabc were collected by centrifugation and resuspended in Middlebrook 7H medium supplemented with 0.2% glycerol, 0.05% Tween 80 and 10% OADC. Frozen aliquots were stored at −70 °C. Representative vials were thawed and the numbers of cfu. viable on Middlebrook 7H10 agar were counted. Single-cell suspensions of mycobacteria were prepared as described previously.23

Reagents

LPS (Escherichia coli serotype 055:B5), ATP disodium salt and DMSO (dimethyl sulfoxide; added to the cultures at 0.1% (v/v) as a solvent control) were from Sigma-Aldrich. For the experiments employing elevated potassium levels (150 mm; Sigma), sodium was replaced by potassium at an equivalent molar concentration.54 BAPTA-AM and z-YVAD-fmk were from Tocris Bioscience (Ellisville, MO, USA). For western blot analysis, Syk (N-19), p62 SQSTM1 (H-290), MyD88 (N-19), LL-37 (H-40) and β-actin (I-19) Abs were from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Human caspase-1 p45 (sc-622) and p10 (sc-515) were purchased from Santa Cruz Biotechnology. Human IL-1β was from Cell Signaling Technology (Danvers, MA, USA). Mouse anti-human dectin-1/CLEC7A mAb was from R&D Systems (Minneapolis, MN, USA) and IC mAb (IgG2a) was purchased from eBioscience (San Diego, CA, USA).

shRNA transduction

Human pLKO.1 lentiviral constructs from Open Biosystems (Huntsville, AL, USA) were used that target NLRP3 (NM_001079821), ASC (NM_013258), LL-37 (NM_004345) and p62 (NM_00142928) were supplied by Sigma-Aldrich as glycerol stocks. Virus production was performed as described previously.59 Briefly, lentiviruses were transfected into HEK 293T cells with pLKO puro.1 or target shRNA plasmids and the packaging plasmids pMDL-RRE (5 μg), pRSV-REV (2.5 μg) and pVSV-g (3 μg). After 48 h, HEK 293T cell supernatants with target-containing virus were collected. The nontarget pLKO.1 shRNA (shNS) or target lentiviral particles were mixed with 8 μg/ml Polybrene (Sigma-Aldrich) in HEKs, according to the manufacturer’s protocol. After 3 days, the transduction efficiency was determined by RT–PCR.

Transfection of siRNA into THP-1 cells

RNA interference experiments were performed as described previously.39 Briefly, the psiRNA–h7SKGFPzeo plasmids for human TLR2 and TLR4 were purchased from InvivoGen (San Diego, CA, USA). Human Syk-targeting siRNA (sc-29501; a mixture of three target-specific 20–25 nt siRNAs) and control nontargeting siRNA (sc-37007) were purchased from Santa Cruz Biotechnology. Cells were transfected with 20 μl of scrambled siRNA or specific siRNA (dectin-1, Syk and MyD88; Santa Cruz Biotechnology) with the transfection reagent Lipofectamine 2000 (Invitrogen), according to the manufacturer’s protocol. Transfected cells were stimulated with Mabc after being harvested for western blot analysis.

RNA extraction and RT–PCR analysis

RNA extraction from monocytes and THP-1 cells was performed as described previously.34,39 Briefly, total RNA was extracted used TRIzol reagent (Invitrogen). PCR was performed with a Peltier Thermal Cycler-200 (Watertown, MA, USA) using 35 cycles with 45 s of annealing, and temperatures as follows: 52 °C for p62 and β-actin, 54 °C for TLR2 and TLR4, 58 °C for ASC and NLRP3, 59 °C for LL-37, 60 °C for dectin-1, 65 °C for DEF4 and 70 °C for IL-1β. The primer pairs are shown in Supplementary Table 1.

Western blot and ELISA analysis

Western blot analysis using whole-cell lysates and cell supernatants was performed as described previously.46,55 Briefly, after stimulation, nuclei and cell debris were removed by centrifugation (20 min; 14 000 g). The post-nuclear supernatants were further cleared by centrifugation at 100 000 g for 1 h. Supernatants were then run on SDS-PAGE and blotted onto PVDF membranes. Blots were probed with the following primary Abs: human PMS (I-19), MyD88 (N-19), LL-37 (H-40) and β-actin (I-19) Abs were from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Human caspase-1 p45 (sc-622) and p10 (sc-515) were purchased from Santa Cruz Biotechnology. Human IL-1β was from Cell Signaling Technology (Danvers, MA, USA). Mouse anti-human dectin-1/CLEC7A mAb was from R&D Systems (Minneapolis, MN, USA) and IC mAb (IgG2a) was purchased from eBioscience (San Diego, CA, USA).
60 min at 4 °C. To concentrate the secreted IL-1β from cell culture media, 10 μl of StrataClean resin (Stratagene, La Jolla, CA, USA) were added to the each sample (200 μl), which were then incubated for 30 min at 4 °C, according to the manufacturer's protocol. After resin-bound protein, samples were washed twice with phosphate-buffered saline by centrifugation, resuspended with 1× sample buffer and analyzed by western blot analysis. Cytokine levels were measured by ELISA (for IL-1β, TNF-α and IL-8 (BD Pharmingen)), as described previously.49

Intracellular Ca2⁺ measurements
Intracellular Ca2⁺ measurements were performed as described previously.56 Briefly, THP-1 cells grown on coverslips were loaded with the Ca2⁺ indicator Fluo-4/AM (10 μM, 30 min for kinetic measurements) in HBSS (Hank's balanced salt solution) buffer, according to the manufacturer's protocol (Molecular Probes, Carlsbad, CA, USA). The cells were washed twice with HBSS buffer and infected with Mabc. For confocal measurements, images were obtained with an LSM510 confocal microscope (Zeiss, Thornwood, NY, USA).

Quantification of mycobacterial growth
Briefly, the capacity of intracellular survival was examined using Mabc-induced human MDMs. After 3 days of culture, cells were trypsinized and lysed with 0.025% Triton X-100 to collect intracellular bacteria. The lysates were then resuspended and sonicated in a preheated 37 °C bath sonicator (Elma, Singen, Germany) for 5 min. Aliquots of the sonicates were serially diluted in 7H9 broth, plated separately on 7H10 agar plates for colony counting in triplicate and incubated at 37 °C with 5% CO2 for 7 days.

Statistical analyses
All data are presented as mean values ± s.d. of independent determinations. For statistical analyses, paired t-tests with Bonferroni adjustment or analysis of variance for multiple comparisons were performed. Differences were considered significant at P<0.05.

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