

Innate immunity holding the flanks until reinforced by adaptive immunity against Mycobacterium tuberculosis infection

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Provisional

1 **Innate immunity holding the flanks until reinforced by adaptive immunity against**
2 ***Mycobacterium tuberculosis* infection**

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27 **Abstract**

28 T cells play a cardinal role in imparting adaptive immunity against *Mycobacterium tuberculosis*
29 (*Mtb*). However, ample time is required before T-cells are able to evoke efficient effector
30 responses in the lung, where the mycobacterium inflicts disease. This delay in T cells priming,
31 which is termed as lag phase, provides sufficient time for *Mtb* to replicate and establish itself
32 within the host. In contrast, innate immunity efficiently curb the growth of *Mtb* during initial
33 phase of infection through several mechanisms. Pathogen recognition by innate cells rapidly
34 triggers a cascade of events, such as apoptosis, autophagy, inflammasome formation and nitric
35 oxide production to kill intracellular pathogens. Furthermore, bactericidal mechanisms such as
36 autophagy and apoptosis, augment the antigen processing and presentation, thereby contributing
37 substantially to the induction of adaptive immunity. This manuscript highlights the role of innate
38 immune mechanisms in restricting the survival of *Mtb* during lag phase. Finally, this article
39 provides new insight for designing immuno-therapies by targeting innate immune mechanisms to
40 achieve optimum immune response to cure TB.

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51 **Introduction**

52 Tuberculosis (TB) continues to affect public health worldwide. About one third of the global
53 population is infected with *Mtb* but only 3%-10% succumb to disease (Barry et al.,
54 2009;Ottenhoff and Kaufmann, 2012). Therefore, greater than >90% of infected population
55 remains asymptomatic, which determines the intricate balance between host immunity and *Mtb*.
56 Understanding the immune response of these asymptomatic individuals can be highly
57 informative and will provide potentially new pathways for the development of anti-TB drugs and
58 vaccines.

59 Over the past several decades, research related to defense against *Mtb* was largely focused on the
60 T cells because of their remarkable ability to generate *Mtb* specific immunity, followed by an
61 enduring memory response to counter subsequent infections (Stenger and Modlin, 1999).
62 Undoubtedly, T cells play a crucial role in protection against *Mtb*. However, recent information
63 has chiseled the long belief that T cells are the sentinels in *Mtb* protection, in part due to the
64 substantial lag period between infection and the establishment of specific T-cell responses
65 (Shaler et al., 2012). Recruitment of dendritic cells (DCs) to the site of infection, followed by
66 their *Mtb* acquisition and transportation towards draining lymph nodes to prime naïve T cells
67 takes 9-11 days after the invasion of the pathogen. Hence, T cell activation occurs after
68 considerable time of infection. This delay between the onset of infection and generation of
69 specific effector T cells provides enough time for *Mtb* to establish an infection [Fig. 1]. Once
70 established, *Mtb* ultimately hampers the antigen processing and priming of naïve T cells (Roberts
71 and Robinson, 2014). Eventually, obstructs the generation and propagation of anti-*Mtb* T cell
72 responses. However, despite the lag phase in T-cell responses, >90% of infected individuals are
73 asymptomatic, raising the possibility of the involvement of other factors in controlling TB.

74 Lurie's fundamental studies with resistant and susceptible inbred rabbits proved that the innate
75 response effectively controls the growth of *Mtb* at early times of infection. After 7 days of
76 inhalation of tubercle bacilli, lungs of susceptible animals showed 20 to 30 fold more bacteria
77 than resistant strain (Lurie, 1964;Arthur M. Dannenberg and Rook, 1994) (van Crevel et al.,
78 2002) [Fig. 1]. Protection during the initial phase of infection clearly indicates that T cells are not
79 at the forefront in controlling the infection, but, rather that components of the innate immune
80 system play a pivotal role in generating efficient immunity against *Mtb*. Therefore, it is important
81 to dissect the mechanisms responsible for curbing the growth of *Mtb* during the lag phase of T
82 cell response. Understanding these mechanisms could pave ways in designing novel therapeutic
83 strategies and vaccines to enhance the immune response more efficiently against *Mtb*.

84 **Why focus more on innate immunity?** In the past, the role of innate immunity was ignored in
85 inducing a protective response against *Mtb*. Recent reports show that the function of innate
86 immunity is even more effective than T cell response against *Mtb* (Fremond et al., 2004;Nicolle
87 et al., 2004). Mice with defective MyD88 signaling show optimal T cell response, yet there is no
88 significant reduction in the lung bacterial burden of *Mtb* challenged mice, compared to wild type
89 (Nicolle et al., 2004). In another study it was shown that MyD88 knockout mice show
90 interferon-(IFN)- γ production in response to mycobacterial antigens but *Mtb* infection became
91 lethal within 4 weeks of post infection with 2 log₁₀ higher CFUs in the lung (Fremond et al.,
92 2004). TLR-2 deficient or TLR-9 KO mice show high levels of IFN- γ and tumor necrosis factor
93 (TNF)- α , with high infiltration of CD4 T cells and CD8 T cells in lungs but these mice succumb
94 to *Mtb* infection (Carlos et al., 2009). These reports indicate that innate immunity has a more
95 profound role than to simply assist adaptive immunity. Moreover, an optimal acquired immune
96 response is not sufficient to compensate for defective innate immunity. Collectively, these

97 studies suggest that it is important to dissect out the functions assisted by innate immunity to
98 induce a protective response against *Mtb*.

99 **Innate mechanisms act and foster T cells to react against *Mtb*.** Based on the current studies,
100 innate immunity has gained much more impetus due to its profound role in early control of *Mtb*
101 infection and in sustaining the T cell response (Sia et al., 2015). Whether or not these
102 mechanisms represent the target to explore in designing new strategies to control *Mtb* needs to be
103 explored. Taking into consideration all these facts, herein we compile the contribution of the
104 bactericidal mechanisms: autophagy, apoptosis, inflammasome formation, and nitric oxide
105 production in limiting the growth of *Mtb* (Table 1). Additionally, we will discuss how innate
106 signaling delivered through pattern recognition receptors (PRRs) such as toll like receptors
107 (TLRs), nucleotide binding oligomerization domain like receptors (NLRs) can augment these
108 mechanisms.

109 **Autophagy.** Autophagy is evolved as a stress response that endows cells with a capability to
110 adjust their biomass and turn over constituents at the time of starvation. It targets the cytoplasmic
111 material, including macromolecules, organelles and cells undergoing unscheduled apoptosis to
112 lysosomes for degradation, thus periodically cleaning their interiors. Furthermore, autophagy has
113 crucial roles in various biological processes, which include aging, development, degenerative
114 diseases, and cancer (Huang and Brumell, 2014; Jiang and Mizushima, 2014). In addition, it also
115 helps in elimination of pathogens, which exploit the cytosolic compartment for their regular life
116 cycle, or those that are evolved with the capability to arrest phago-lysosome biogenesis
117 (Flannagan et al., 2009). Autophagy is initiated with the sequestering of pathogens intracellularly
118 to form a double membrane envelope that is known as autophagosome. Autophagosome fuses
119 with lysosomes to form autolysosome to degrade pathogens. Thereby, autophagy facilitates the

120 trafficking of mycobacteria to the lysosome for degradation (Gutierrez et al., 2004; Deretic et al.,
121 2009). Similar results are reported in the case of *Bacillus Calmette–Guerin* (BCG). In addition,
122 autophagy also transports a large proportion of ubiquitinated proteins to lysosomes and augments
123 the bactericidal capacity of lysosomal fraction (Alonso et al., 2007). *Mtb* escapes the immune
124 mechanism by neutralizing the acidification of phagosomes (Deretic et al., 2006; Russell, 2011).
125 Autophagy overcomes this *Mtb* evasion strategy by targeting phagosomes containing bacterium
126 to lysosomes (Jo, 2013). Thus, autophagy provides an additional barrier to neutralize an attempt
127 made by the mycobacterium to manipulate phagosomes maturation.

128 Noteworthy, autophagy processes bacteria for degradation within early hours of infection, as
129 evidenced by conversion of autophagy marker LC3-I to LC3-II in *Mtb* infected DCs and Mφ
130 (Khan et al., 2016). This experiment categorically indicates that autophagy guards the host
131 against *Mtb* during the initial phase of infection. Importantly, animals with defective autophagy
132 showed increase in the bacterial burden in lungs of *Mtb* challenged animals despite of
133 predominance of Th17 immunity (Castillo et al., 2012). Furthermore, these animals showed
134 remarkable gross tubercle lesions, in contrast to the smaller infected foci in the lungs of control
135 animals. It signifies that autophagy also aid in preventing excessive inflammatory reactions in
136 the host.

137 Currently, the only available vaccine for TB is BCG. Nonetheless, BCG has failed to reduce the
138 global TB burden. Interestingly, one of the factors associated with the failure of BCG in TB-
139 endemic areas is BCG ability to hamper the fusion of phagosome with lysosomes (Soualhine et
140 al., 2007;Sun et al., 2007;Gowthaman et al., 2012). This interference in the antigen processing
141 and presentation to T cells, results in defective T cell response. Since autophagy can overcome
142 the problem of phago-lysosome biogenesis, targeting autophagy can substantially contribute to

143 improve efficacy of BCG vaccine (Jagannath et al., 2009). It has been shown in a very elegant
144 study that mice immunized with rapamycin-treated DCs infected with BCG showed enhanced
145 Th1 cells mediated protection when challenged with virulent *Mtb*. Rapamycin induced
146 enhancement in antigen presentation was attenuated when autophagy was suppressed by 3-
147 methyladenine or by small interfering RNA against beclin-1 (Jagannath et al., 2009). Targeting
148 autophagy may open new avenues in boosting the efficacy of BCG or immune response against
149 *Mtb*. Dissecting the host factors that regulates autophagy can help in restricting the growth of
150 *Mtb* and simultaneously improving the processing and presentation of antigen and enhancing T
151 cell immunity.

152 Autophagy can be induced due to starvation, treatment with IFN- γ or rapamycin (Bento et al.,
153 2015). Additionally, triggering through PRRs has direct correlation with the induction of
154 autophagy. Signaling through TLR-4, TLR-3, TLR-7 and NOD-2 receptor can potently induce
155 autophagy (Delgado et al., 2008;Delgado and Deretic, 2009;Cooney et al., 2010;Yuk et al.,
156 2012). TLR-7 triggering enables M ϕ s to reduce the survival of *Mtb* via induction of autophagy
157 (Delgado et al., 2008). It was further supported through suppression of autophagy by knocking
158 down beclin and Atg5 through siRNA. Innate triggering through NOD-1 and NOD-2 enhances
159 autophagy induction and reduces the bacterial burden within 6h of infection (Travassos et al.,
160 2010). Murine immunity related to guanosine tri-phosphate induces autophagy and generate
161 large autolysosomal organelles, as a mechanism for the elimination of intracellular *Mtb*.
162 Furthermore, human Irgm1 ortholog (IRGM) augments autophagy and reduces intracellular
163 bacillary load (Singh et al., 2006). Many evidences have been documented to show the potential
164 for autophagy based therapies to target *Mtb* (Bento et al., 2015). Similarly, exploring innate
165 receptors to augment autophagy could also be one of the strategies to boost the host immunity

166 against *Mtb*. Moreover, it would be beneficial in overcoming the failure associated with BCG
167 vaccine.

168 **Reactive nitrogen and oxygen intermediates.** Nitric oxide (NO) and reactive nitrogen
169 intermediates (RNI) are considered potent antimicrobial agents acquired by innate cells. M ϕ s are
170 the major producer of NO. It is released by inducible nitric oxide synthase (iNOs), a heme-
171 protein that catalyses the oxidation of L-arginine to NO and citrulline. NO production is a critical
172 defense mechanism in determining the outcome of TB infection, since it reduces the survival of
173 *Mtb* (Rich et al., 1997). This information was corroborated with the result observed in mouse
174 model where abrogation of iNOs activity produces dramatic increase in microbial burden
175 (MacMicking et al., 1997). Further, disruption of iNOs gene expression results in a high rate of
176 *Mtb* dissemination and mortality. Stimulation of innate molecules, such as TLRs or NODs
177 trigger the expression of iNOs, which ultimately kills the *Mtb*, as evidenced by colony forming
178 units (CFU) assay (Chan et al., 2001). In the mouse model of TB, NO secretion is well known to
179 be an antimicrobial defense mechanism. However, its role in humans is still controversial.
180 Forthcoming evidences indicate that human M ϕ s and alveolar epithelial cells upon infection with
181 *Mtb* secrete NO to inhibit the intracellular growth of *Mtb*. Additionally, iNOs and markers
182 associated with NO are highly expressed in the M ϕ s obtained from broncho-alveolar lavage of
183 TB patients and not healthy individuals (Nicholson et al., 1996). Interestingly, patients infected
184 with multidrug resistant (MDR) strain of *Mtb* produce less NO (Sharma et al., 2004). More
185 startling observation came from the report that chemotherapy appears to cure *Mtb* in immune-
186 competent mice but fails to do so in NOS2-deficient animals (Nathan and Shiloh, 2000). It
187 concludes that bactericidal drug uses NO pathway for efficient killing of *Mtb* (Ciccone et al.,
188 2003).

189 IFN- γ induces the production of NO (Flesch and Kaufmann, 1991). After several days of *Mtb*
190 infection, T cells produce IFN- γ . However, production and action of NO is observed within 3h
191 and it persists for few days in the circulation (Akaki et al., 1997; Rich et al., 1997). Early
192 production of NO signifies that IFN- γ , which is a potential stimulator for NO release, is not
193 being released by T cells, but instead by the cells of innate immunity such as natural killer (NK)
194 cells and $\gamma\delta$ T cells (Sada-Ovalle et al., 2008). It suggests that innate immune cells are
195 responsible for early release of NO to restrict the growth of *Mtb* during the initial period of
196 infection. NO is also reported to regulate the synthesis and release of several pro-inflammatory
197 cytokines including IL-1 β , TNF- α and IL-8, which subsequently affect the production of NO in
198 the feedback loop (Kuo et al., 2000). Microbicidal activity of M ϕ s is also associated with
199 reactive oxygen intermediates (ROI). However, their role in constraining the growth of *Mtb* is
200 not highly significant (Chan et al., 1992; Akaki et al., 1997; Lamichhane, 2011a;b). Probable
201 reason documented is that although ROI appear immediately upon *Mtb* infection but M ϕ s ceases
202 to produce it within 2h of infection. Further, ROI also have shorter half-lives.

203 In essence, the NO kills *Mtb* and as well as its augment of the host immunity against the
204 pathogen is well documented in the literature (Chan et al., 2001). It is important to mention that
205 TLRs signaling contributes substantially in release of NO (West et al., 2011). 1, 25-
206 dihydroxyvitamin D is a potent inducer of NO and suppresses the growth of *Mtb* (Rockett et al.,
207 1998). TLRs triggering enhances the bactericidal activity of M ϕ s by upregulating the expression
208 of the vitamin D receptor and by inducing the enzyme that catalyzes the conversion of 25-
209 dihydroxyvitamin D3 to active 1, 25-dihydroxyvitamin D leading to the induction of
210 antimicrobial peptide cathelicidin (Liu et al., 2006). Hence, it may be considered as a critical
211 molecule to consider while designing new therapeutic strategies to treat TB. TLR ligands are

212 known to induce NO production in antigen presenting cells (APCs) and NO restricts the growth
213 of *Mtb* (Khan et al., 2015).

214 **Apoptosis.** Apoptosis is commonly known as ‘programmed cell death (PCD)’. It is a
215 phenomenon that occurs when a cell committing suicide confines its cytoplasmic content within
216 membrane bound vesicles named as apoptotic bodies. These membrane bound vesicles express
217 molecules known as ‘eat me or find me’ signals. ‘Eat me’ signals helps in the recognition of
218 these unwanted moieties by phagocytic cells (Behar et al., 2011). Furthermore, phagocytic cells
219 remove them through a mechanism known as efferocytosis; the process known to engulf and
220 remove apoptotic cells. Failure in efferocytosis results in the disintegration of apoptotic bodies
221 and release of intracellular contents. This causes inflammation that is known as secondary
222 necrosis (Martin et al., 2012). Importantly, apoptosis makes a crucial contribution to the host
223 immune response and determines the outcome of infection. It abolishes the protected
224 intracellular niche favoring the replication of *Mtb*, thus forcing the bacteria to search for a new
225 habitat. The caspase family of serine proteases is the central molecules responsible for the
226 execution of apoptosis. Apoptosis is classically induced by three pathways. First is through
227 ligation or oligomerization of tumor necrosis factor receptor (TNFR) family. Ligation of cell
228 surface receptor such as TNFR or Fas, results in the subsequent activation of caspases and the
229 induction of apoptotic vesicles. Intrinsic apoptosis occurs in response to oxidative stress, nutrient
230 starvation or intracellular stress, which changes the mitochondrial membrane permeability. It
231 results in the translocation of cytochrome c from mitochondria to the cytoplasm, leading to the
232 activation of caspases. The third pathway is mediated by granzyme B released from cytolytic T
233 cells and NK cells.

234 *Mtb* induces apoptosis through the classical extrinsic pathway. Encounter of *Mtb* with innate
235 cells such as DCs and M ϕ s induces the release of TNF- α and triggers apoptosis. Apoptosis limits
236 the replication of *Mtb* by sequestering bacilli in apoptotic vesicles and by activating nearby
237 uninfected M ϕ s. This phenomenon has been demonstrated through a classical experiment in
238 which uninfected autologous M ϕ s were cultured with apoptotic or necrotic or non-apoptotic
239 infected M ϕ s. Interestingly, significant inhibition in the growth of *Mtb* was seen when apoptotic
240 cells were cultured with uninfected M ϕ s. In coculture experiments, elimination of *Mtb* was
241 anticipated through efferocytosis. Later, antimicrobial effect enacted by naïve M ϕ s was shown to
242 be contact independent (Hartman and Kornfeld, 2011). Interleukin-1 signaling in naïve M ϕ s
243 mediates the cross-talk with infected-M ϕ s. It exhibits NO-dependent antimicrobial activity
244 against bacilli in autolysosomes of heavily infected M ϕ s (Hartman and Kornfeld, 2011).
245 Noteworthy, the discrepancy occurs in the induction of apoptosis by avirulent *versus* virulent
246 *Mtb* (Chen et al., 2006). Multiple reports indicate that virulent *Mtb* induces necrosis to avoid host
247 defensive strategies, whereas attenuated strain is associated with apoptosis (Chen et al., 2006;
248 Divangahi et al., 2009). Despite comparable amount of TNF- α , cells infected with avirulent
249 strain are more susceptible to apoptosis. It was revealed that difference in level of apoptosis
250 between *Mtb* strains is due to an evasion strategy used by the virulent strain of *Mtb*. Cells
251 infected with virulent *Mtb* secrete more IL-10, which induces the release of TNFR-2. Soluble
252 TNFR-2 forms a complex with TNF- α and downregulates the TNF- α induced apoptosis
253 (Balcewicz-Sablinska et al., 1998). Furthermore, it has been demonstrated by Annexin V binding
254 and intracellular caspase staining that early secretory antigen target (ESAT)-6 of *Mtb* induces
255 apoptosis in human M ϕ s (Choi et al., 2010). Additionally, the expression profile of apoptotic
256 genes shows up-regulation of anti-apoptotic genes in virulent *Mtb* infected M ϕ s.

257 In addition to the restriction of the *Mtb* growth during early phase of infection, apoptosis has a
258 considerable role in the induction of the acquired cellular immune response (Winau et al., 2006).
259 Both CD4 T cells and CD8 T cells are well documented in immunity against *Mtb*. However, the
260 mechanism underlying the presentation of antigens to CD8 T cells in context with MHC-I
261 molecules remains enigmatic. Recently, it has been shown that apoptosis of infected Mφs
262 facilitates the release of mycobacterial antigens in apoptotic vesicles, thereby allowing their
263 access to bystander APCs to present antigen to CD8 T cells. Inhibition of apoptotic blebbing
264 using caspase inhibitors, hampers the CD8 T cell response (Winau et al., 2006). Therefore, it
265 may be concluded that triggering of apoptosis can efficiently control the *Mtb* growth at early
266 time points; and at later stages it potentially contributes in the generation of antigen specific CD8
267 T cells.

268 Neutrophils are important cells of innate immunity. They play a significant role in imparting
269 protection to *Mtb* (Andersson et al., 2014). These are the first cells to be recruited at the site of
270 infection. Neutrophils phagocytose *Mtb*. Furthermore, *Mtb* infected neutrophils undergo
271 apoptosis and are phagocytosed by Mφs. These Mφs then release TNF-α to form granulomas and
272 control acute *Mtb* infection (Perskvist et al., 2002). Further, inhibition of apoptosis in neutrophils
273 delays the priming of CD4 T cells. Hence it implies that apoptosis plays a decisive role in
274 controlling *Mtb* infection by activating innate as well as adaptive immunity. TLRs induced
275 apoptosis such as TLR-3, 4 has been explored for cancer therapy (Salaun et al., 2007).
276 Interestingly, TLRs show enough potential for triggering apoptosis in *Mtb* infected cells. LpqH,
277 a 19 kDa and 38 kDa lipoprotein of *Mtb* induces the Mφ cell death in TLR-2 dependent manner
278 (Ciaramella et al., 2000; Sanchez et al., 2012). 38 kDa lipoprotein of *Mtb* elicits the TNF-α
279 release in TLR-2 dependent manner and induces apoptosis in infected Mφ (Sanchez et al., 2009).

280 Apoptosis has been shown to improve the efficacy of BCG vaccine. Deletion of the secA2 gene
281 of *Mtb*, which encodes a component of a virulence-associated bacterial protein, triggers the
282 apoptosis of infected cells and enhances the priming of antigen specific CD8 T cells. Vaccination
283 with secA2 deleted *Mtb* mutant induces better protection than BCG against *Mtb* (Boom,
284 2007;Hinchey et al., 2007). rBCG strain that secretes listeriolysin of *Lysteria monocytogens*
285 induces more efficacious protection than BCG against *Mtb* by facilitating the cross priming by
286 inducing apoptosis (Grode et al., 2005). This evidence indicates that targeting apoptosis could be
287 one of the potential strategies to prevent TB.

288 Although apoptosis is the well-studied PCD, but it is not the only mechanism responsible for this
289 process. A new form of non-apoptotic PCD has been termed as paraptosis. Insulin like growth
290 factor I receptor has been identified as a molecule involved in inducing paraptosis. It is
291 characterized by cytoplasmic vacuolation, along with mitochondrial swelling, lack of apoptotic
292 morphology, caspase activation and inhibition by caspase inhibitors (Sperandio et al., 2004). A
293 few reports suggest that this form of cell death is driven by an alternative caspase-9 activity that
294 is Apaf-1-independent (Sperandio et al., 2000). Since, paraptosis follows the pathway different
295 from apoptosis, it could be a novel therapeutic target to kill pathogens that inhibits apoptosis.
296 Little is known about the effect of paraptosis on the immune system and moieties involved in it.
297 A few apoptotic inducers have been shown to elicit paraptosis (Amarante-Mendes et al., 1998).
298 Nothing is known about its role in TB. In the future, it may be an interesting line of investigation
299 to understand the contribution of paraptosis in limiting the *Mtb* growth.

300 **Inflamasome.** *Mtb* activates the cascade of events mediating the release of an array of pro-
301 inflammatory cytokines such as IL-6, IL-12 and TNF- α that play a defensive role in eliciting
302 innate immunity (Cooper et al., 2011). Similarly, IL-1 β and IL-18 have an influential role in

303 imparting protection to *Mtb*. IL-18 enhances the production of IFN- γ and its abrogation results in
304 less IFN- γ release and impaired NK cell function (Kawakami et al., 2000). Simultaneously, IL-
305 1R1-deficient mice show 2-log increase in bacterial load in the lung and necrotic pneumonia
306 within 4 wks of *Mtb* exposure. It is notable to mention that cell mediated immunity (CMI), which
307 is considered the hallmark of protection against *Mtb* is not sufficient in restricting bacterial
308 burden in IL-1R deficient mice, despite efficient pulmonary CD4 T cells and CD8 T cell
309 responses (Fremond et al., 2007).

310 Unlike other proinflammatory cytokines, IL-1 β and IL-18 are synthesized as precursors known
311 as pro-IL-1 β and pro-IL-18 (Sansonetti et al., 2000). Multiple signaling pathways triggered
312 through TLRs and cytokines result in the transcription of pro-IL-1 β and pro-IL-18. However,
313 their maturation requires processing by active caspases. Distinct caspases regulate the apoptosis
314 and maturation of IL-1 β and IL-18. Caspase-1 regulates the maturation of IL-1 β and IL-18
315 (Dinarello, 2006). Importantly, release of IL-1 β and IL-18 is highly regulated phenomenon,
316 which is dependent on the activation of caspase-1 and its homolog by multimeric protein
317 complex termed as inflammasomes (Vladimer et al., 2013). These complexes are critical in the
318 proteolytic processing of pro-IL-1 β and pro-IL-18 into their active form (Netea et al.,
319 2010;Briken et al., 2013). The inflammasome is classically composed of NOD like receptors
320 (NLRs), the adaptor molecule PYCARD/ASC, and pro-caspase-1, which when proteolyzed to
321 caspase-1 provides the enzymatic activity of the inflammasome. Pro-caspase-1 forms the core of
322 the inflammasome. However, the constitution of NLRs within the inflammasome varies
323 according to the type of pathogen involved. The NLR family members NALP3, NAIP5 or IPAF
324 and the adaptor apoptosis speck-like protein (ASC) are involved in caspase-1 activation.
325 Inflammasome plays an important role in host defense against *Mtb*, since mice deficient in IL-1

receptor (IL-1RI), IL-1 β or IL-18 are more susceptible to infection with *Mtb*. Furthermore, a defect in ASC adaptor protein shows the exacerbation of disease without restricting the *Mtb* growth. Early secreted antigenic target protein 6 kilodalton secretion system (ESX)-1 encoded in RD-1 region of *Mtb* promotes the release of IL-1 β by inflammasome activation. ESX-1 mediated inflammasome formation depends on host NLRP-3 and ASC protein (Mishra et al., 2010). RD-1 deficient *Mtb* fails to induce a strong activation of caspase-1 resulting in inefficient secretion of IL-1 β and IL-18. This observation signifies that the failure of BCG to mount optimal protection against *Mtb* is due to absence of RD-1 dependent induction of IL-1 β and IL-18 (Kurenuma et al., 2009). Interestingly, treatment with exogenous IL-18 reduces the bacterial load in mice. Recently, viral and bacterial RNA have been shown to trigger NLRP3 and activate inflammasome (Mitoma et al., 2013). It suggests that prophylactic strategies employing recombinant BCG expressing innate ligands, which are efficient in inducing inflammasome formation, can boost its protective efficacy against TB. *Mtb* genes Rv0198c (zmp1), plays a critical role in preventing caspase-1-dependent activation and secretion of IL-1 β . zmp1-deleted *Mtb* triggered activation of the inflammasome, resulting in increased release of IL-1 β , enhanced maturation of *Mtb* containing phagosomes, improved mycobacterial clearance by macrophages, and reduction in bacterial load in the lungs of *Mtb* aerosol-infected mice (Master et al., 2008). Zmp1 is an important virulence determinant and represents a potentially useful drug target. Furthermore, it has been shown that binding of vitamin D induces IL-1 β secretion and prevent infection. This information supports the idea of exploiting vitamin D in clinical trials against *Mtb* (Verway et al., 2013).

Inflammasomes are also reported to play an important part in amplifying the adaptive immune response. Importantly, inflammasome processed IL-1 β promotes the differentiation of naïve CD4

349 T cells to Th17 subtype. It synergizes with IL-6 and promotes Th17 cell development *via* up-
350 regulation of key cytokine IL-17, transcription factors, IRF4 and ROR γ t. Furthermore, IL-1 β can
351 coordinate with IL-6 and IL-23 in the absence of TGF- β signaling to induce pathogenic Th17
352 cells (Ghoreschi et al., 2010). In addition to Th1 cells, Th17 cells also play a cardinal role in
353 generating anti-*Mtb* response. IL-17 induces the expression of chemokines that results in the
354 recruitment of various cells to site of infection. Furthermore, memory Th17 cells promote rapid
355 migration of Th1 cells by enhancing the expression of chemokines (Khader and Cooper, 2008).
356 Toll-like receptors and NOD-2 expressed on antigen presenting cells are responsible for the
357 induction and release of cytokines like IL-6, TGF- β , and IL-12 that are responsible for the
358 differentiation of Th17 cells and Th1 cells, respectively (Khan et al., 2016). Hence
359 immunotherapies involving agonists of innate immunity can be explored in the generation of
360 protective immunity against *Mtb* (Chodisetti et al., 2015). The above showcased points indicate
361 that innate immunity efficiently controls the *Mtb* growth during early phase of infection.
362 Moreover, it creates a platform for adaptive immunity.

363 Conclusion

364 Continuous efforts are undertaken to generate an effective vaccine against TB. However, a
365 possible candidate that can achieve the WHO-STOP-TB program has not yet been formulated.
366 Eleven candidate vaccines are currently in clinical trials. Failure of BCG to protect against *Mtb*
367 warrants a serious attempt to reinvigorate BCG potency for inducing optimal immune response
368 (Singh et al., 2010; Gowthaman et al., 2011; Gowthaman et al., 2012).

369 Recently, innate immunity has emerged as a cornerstone in limiting the growth of *Mtb* (Fremond
370 et al., 2004; Nicolle et al., 2004; Carlos et al., 2009). Innate immunity not only initiates series of
371 events to assist adaptive immunity but also restricts the growth of TB bacilli at the initial phase

372 of infection. Nonetheless, failure of innate killing mechanisms results in unobstructed growth of
373 *Mtb* and provides enough opportunity for the pathogen to breach the barrier of the immune
374 system. It indicates that targeting innate immunity is a judicious approach to consider while
375 designing vaccines or therapeutics. Inefficiency of innate immunity provides an opportunity for
376 unimpeded *Mtb* growth. Later, the *Mtb* conquers the adaptive immunity. Adequate innate
377 immunity is capable of restricting the growth of *Mtb* during the “lag phase” of T cell response.
378 Limiting the growth of *Mtb* during the initial phase of infection provides enough time for T cells
379 to reach the site of infection and curtail *Mtb* replication. However, impaired innate immunity is
380 incompetent in curbing the proliferation of *Mtb*. It results in unhindered growth of *Mtb*, which
381 ultimately interferes in the activation of adaptive immunity. Biological therapies involving innate
382 ligands for TLRs and NLRs will benefit the quest for novel treatment modalities for TB. We
383 speculate that mycobacterial vaccines engineered with ligands for PRRs may enhance the
384 potency of innate immunity to limit the *Mtb* growth and sustain the adaptive arm of immunity.

385

386 **Abbreviations**

387 *Mycobacterium tuberculosis*: *Mtb*; dendritic cells: DCs; tuberculosis: TB; *Bacillus Calmette–*
388 *Guerin*: BCG; programmed cell death: PCD

389

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394

395 **Declaration of interest**

396 Authors declare no conflict of interest.

397

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654 **Figure Legends**

655 **Fig. 1. Innate immunity restricts the bacterial burden during lag phase of T cell response.**
656 Initiation of T cell response occurs after 9-11 days of *Mtb* infection and peaks at 20-25 days (-).
657 Delay in the duration for the generation of effective T cell response is considered as its “lag
658 phase”. Susceptible strain (-) of rabbit shows high bacterial burden during lag phase of T cell
659 response; whereas resistant strain (-) signifies lesser bacterial burden. However, after initiation of
660 T cell response, both the strains restrict *Mtb* growth. It indicates that lesser bacterial burden
661 during lag phase of T cell response in resistant strain of rabbit is due to involvement of innate
662 immunity.

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664 **Table 1**

Mechanism	Function	References
Autophagy	Provide the alternative route for antigen processing and presentation. Target cytosolic antigen to lysosome for degradation. Overcome the evasion strategy of <i>Mtb</i> to inhibit phagolysosome biogenesis.	(Jagannath et al., 2009;Cooney et al., 2010)
Apoptosis	Facilitates the presentation of antigen to CD8T cells. Restrict the bacterial burden.	(Winau et al., 2006;Andersson et al., 2014)
Inflammasome	Involved in maturation of IL-1 β and IL-18.	(Fremond et al., 2007)
Nitric oxide	Intracellular killing of pathogen. Regulate IL-1 β secretion to control inflammation.	(Flesch and Kaufmann, 1991;Chan et al., 1992;Nicholson et al., 1996;Akaki et al., 1997;MacMicking et al., 1997;Rich et al., 1997;Kuo et al., 2000;Nathan and Shiloh, 2000;Chan et al., 2001;Ciccone et al., 2003;Sharma et al., 2004;Lamichhane, 2011a)

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Figure 1.TIF

Fig. 1

