

# Induction and Function of Type I IFNs During Chlamydial Infection

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## Introduction

*Chlamydia trachomatis* infection is the leading sexually transmitted bacterial infection (STI) in the US, as reported by CDC. The global burden of chlamydial infection is likely higher than that reported for STI, as ocular trachoma caused by chlamydiae continues to be the leading cause of preventable blindness in the world [1]. *Chlamydia* spp. also cause significant disease in livestock. In women, *C. trachomatis* is a major cause of pelvic inflammatory disease, ectopic pregnancy, and infertility [2]. Chlamydial infections can be self-limiting, providing evidence for the development of protective immune responses [3, 4]. However, infection induces mostly short-term immunity that is strain (serovar) specific, so the risk of re-infection is high, and carries an increased risk of tissue damaging effects [5]. Human epidemiologic studies also indicate increased risk of disease with repeated infection [6, 7]. Consequently, a great deal of research has focused on understanding chlamydial biology and the immune responses to chlamydial infection, with an obvious goal to develop a vaccine that will induce protective responses to *Chlamydia* while avoiding responses that lead to pathology. In this chapter, we will focus on one such innate immune response, the type I IFNs in chlamydial pathogenesis, with emphasis on their role during infection and the mechanism of induction during chlamydial infection.

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## The Pathogen and Pathogenesis

Gram-negative *Chlamydia* sp. are obligate intracellular pathogens with a relatively small genome (1–1.3 Mbp) and a unique developmental cycle [8, 9]. The first step in the intracellular chlamydial infection is attachment of the infectious and metabolically inactive elementary body (EB) to the host cell surface. Once the EB enters the cell by endocytosis, it modifies the vacuole to inhibit phagolysosome fusion, and remains confined in a membrane bound vacuole, termed the “inclusion,” during its entire developmental cycle [10–12]. Inside the early inclusion the EB transforms into the metabolically active reticulate body (RB) form by a process that involves DNA de-condensation [13] and reductive cleavage of the outer membrane protein complex [14, 15]. Rather than being strictly non-fusogenic with the host vesicular trafficking pathways, the chlamydial inclusion selectively fuses with sphingomyelin containing exocytic vesicles en route to the plasma membrane from the Golgi [16, 17]. Inclusion formation and acquisition of sphingomyelin are initiated very early in the cycle [17], a phenomenon driven by early chlamydial protein synthesis [18]. Inside the inclusion, the RB multiplies by binary fission [10] and the inclusion expands to occupy significant parts of the cytosol during *C. trachomatis* infection. Specific molecular triggers generated in the RBs likely due to its local environment, initiate the conversion of RBs to EBs towards the latter part of the chlamydial developmental cycle, a process that occurs asynchronously. Eventually, by multiple exit mechanisms [19], the infected cells are lysed and the released EBs go on to infect neighboring cells. The sequential conversion from the specialized EB to RB and then back to EB is a unique feature of chlamydial biology.

Unlike several facultative intracellular pathogens, chlamydiae are not equipped with toxins that damage the host cells. *C. trachomatis* strictly infects mucosal epithelial cells during a genital infection and conjunctival cells during an ocular infection. The host response to *C. trachomatis* is initiated by infected epithelial cells [20] and sustained by professional inflammatory cells and neighboring uninfected cells. Chlamydial ligands recognized by surface and intracellular pathogen recognition receptors (PRRs) initiate chemokine and cytokine production as early as 3 h post infection in vivo, suggesting that entry of viable chlamydiae into host cells is sufficient to induce a response [21]. Using the mouse model of chlamydial genital tract infection [22], it has been shown that inflammatory responses are a major determining factor in oviduct pathology. Following bacterial ascension to the oviducts, infected epithelial cells respond to bacterial signals by producing cytokines and chemokines [20] that act locally to recruit PMNs and other immune cells [21]. PMNs are partially protective in the cervix and uterus because they restrict on-going chlamydial replication, amplify cytokine signaling and reduce pathogen load by attacking infected cells [23]. However, PMN recruitment to the oviducts is excessive and prolonged, leading to distal blockage and formation of hydrosalpinx or salpingitis [21, 23–26]. The contribution of innate immune pathways, such as TLR2, IL-1R, IFNAR, TNFR in PMN recruitment and oviduct pathology has been demonstrated using gene knockout mice [27–33]. On the other

hand, CD4<sup>+</sup> Th1 cells that produce a predominant IFN- $\gamma$  response are critical to the control of chlamydial genital and ocular infection, and enhanced Th1 immune responses correlate with protection from infection and disease in both animal models and humans [34–38].

## Induction of Type I IFNs During Chlamydial Infection and Its Biological Implications

Induction of type I IFNs has been observed in multiple cell types including oviduct epithelial cells [39], macrophages [40, 41], fibroblasts (McCoy cells) [42], and mouse DC [43] infected in vitro with multiple chlamydial strains. Therefore, the ability to induce IFN- $\beta$  in response to intracellular *C. trachomatis* infection appears relatively conserved. *C. pneumoniae*, on the other hand does not induce significant levels of IFN- $\beta$  expression in epithelial cells, which could be a result of its ability to degrade TRAF3 [44].

Type I IFNs are largely inhibitory to chlamydial growth during in vitro infections. Early studies showed significant inhibition of *C. trachomatis* infectivity in HeLa cells treated with different isotypes of IFN- $\alpha$  [45]. IFN- $\beta$  treatment of macrophages treated with LPS also resulted in significant killing of *C. psittaci*, as observed with IFN- $\gamma$ -treatment and this was attributed to activation of indoleamine dioxygenase (IDO) activity. IDO decyclizes tryptophan to *N*-formyl kynureine resulting in reduction in tryptophan pool in the cells affecting chlamydial growth [46]. Further, inhibition of chlamydial growth by TNF $\alpha$  was shown to be partly mediated through an autocrine function of IFN- $\beta$  enhancing the activity of IDO and could be blocked by tryptophan [47]. Treatment of murine fibroblasts (L cells) with type I IFNs was also shown to significantly reduce the yield of *C. trachomatis* LGV biovar [48]. Besides its role in chlamydial killing, IFN- $\beta$  was shown to contribute to IFN- $\gamma$  expression and in induction of CXCL10 in mouse macrophages infected with *C. pneumoniae* [40] and *C. muridarum* [41], respectively.

The protective effect of IFN- $\alpha/\beta$  observed in vitro was not recapitulated during in vivo *C. muridarum* infection, both in the lungs and genital tract. In the lung infection model, *Ifnar*<sup>-/-</sup> mice showed less bacterial burden, weight loss, and less pathology in comparison to control mice, which was attributed to lower macrophage apoptosis in the absence of IFNAR signaling [49]. During genital *C. muridarum* infection, *Ifnar*<sup>-/-</sup> mice displayed a slightly enhanced clearance of infection and significantly reduced oviduct pathology [28]. The improved bacterial clearance in *Ifnar*<sup>-/-</sup> mice was associated with an increase in antigen-specific T cells in the iliac nodes, enhanced CD4<sup>+</sup> T cell recruitment to the genital tract and an increased level of the IFN- $\gamma$ -inducible protein, CXCL9 in genital secretion. A similar outcome of overall enhanced infection clearance and reduced pathology was observed in genital chlamydial infection during IFN- $\beta$  neutralization in wild-type mice [30]. However, in this study a slight increase in infectious burden was observed at day 4 post infection during IFN- $\beta$  depletion, which was not sustained and the IFN- $\beta$  depleted mice went on to clear infection at a

faster rate than the mice receiving control sera. A similar outcome of increased infection load at day 4 post infection, which was not sustained was also observed in mice deficient for the transcription factor IRF3, which is essential for IFN- $\beta$  induction. These data suggest that IFN- $\beta$  likely has an anti-chlamydial activity early in infection, but its negative impact on the inflammatory cells and T cells is not protective to the host. Indeed, the T cells from the iliac lymph nodes of *Irf3*<sup>-/-</sup> and *Ifnar*<sup>-/-</sup> mice displayed an enhanced antigen-specific T cell response. IRF3 KO mice also developed significant uterine pathology unlike *Ifnar*<sup>-/-</sup> or IFN- $\beta$  depleted mice, suggesting that IRF3 could play an IFNAR/IFN- $\beta$  independent role in uterine horn protection [30].

Recent discovery of a new family member of type I IFNs, IFN epsilon (IFN- $\epsilon$ ), has generated interest due to its exclusive expression in the mouse and human genital tract [50, 51]. *Ifne*<sup>-/-</sup> mice were shown to have slightly enhanced chlamydial infection, suggesting a protective role in infection. IFN- $\epsilon$  signals by the same receptor, IFNAR used by IFN- $\alpha/\beta$ . Possible explanations for the discordant results during infection between *Ifne*<sup>-/-</sup> and *Ifnar*<sup>-/-</sup> could be a direct role for IFN- $\epsilon$  in chlamydial killing, its regulation by sex hormones and/or a constitutive role in imparting resistance to genital tract epithelia in an IFNAR-independent manner.

To understand the mechanism behind the conflicting role of type I IFN in vitro and in vivo, the pleiotropic immune functions of this cytokine in vivo needs further understanding. Type I IFNs are a potent regulator of adaptive immunity, affecting multiple cell types, including macrophages, lymphocytes, and DCs. IFN $\alpha/\beta$  induces the expression of several interferon response genes (IRG), which are important for Th1 maturation [52]. Type I IFNs have also been implicated in the generation of cytotoxic T cells and promotion of in vivo T cell proliferation [53] and T cell survival [54]. However, type I IFNs are also known to inhibit IFN $\gamma$ -induced MHC class II expression [55–57] a function that contradicts its Th1 stimulatory role. Type I IFNs have also been shown to inhibit maturation and activation of mouse Langerhans cells [58]. IFN- $\beta$  has been reported to augment [59] or downregulate IL-12 and CD40 expression in DC [60]. Further, therapeutic administration of IFN- $\beta$  in multiple sclerosis patients led to inhibition of IL-12, augmentation of IL-10 production [61] and inhibition of IL-1 $\beta$  production [62]. The paradoxical effect of IFN- $\beta$  on the expression of Th1-type immune responses partly depends on the timing of DC exposure (during maturation vs. mature) to IFN $\beta$  [52]. Type I IFNs are also pro-apoptotic and induce the expression of a number of pro-apoptotic genes, which could play a major role in pathological outcomes during infection. Overall, the detrimental effect of IFN- $\beta$  during chlamydial infection is a likely result of inhibition of Th1-response, a reduction in IFN $\gamma$  responsiveness, and induction of an apoptotic response. These results have been largely inferred from gene knockout mice studies and antibody depletion studies. It is possible that the pathological outcome could be different if the mice were treated with recombinant IFN- $\beta$ . Treatment of mice with recombinant IFN- $\beta$  has been shown to downregulate IL-1 $\beta$  levels significantly at multiple steps [62]. Since IL-1 signaling is a major player in oviduct pathology during genital chlamydial infection, this may be protective to the oviducts during infection [29]. Therefore, the overall effects of IFN- $\beta$  during chlamydial infection in vivo is likely determined by its levels in the local tissue, and assigning a beneficial or detrimental role to it would be contextual.

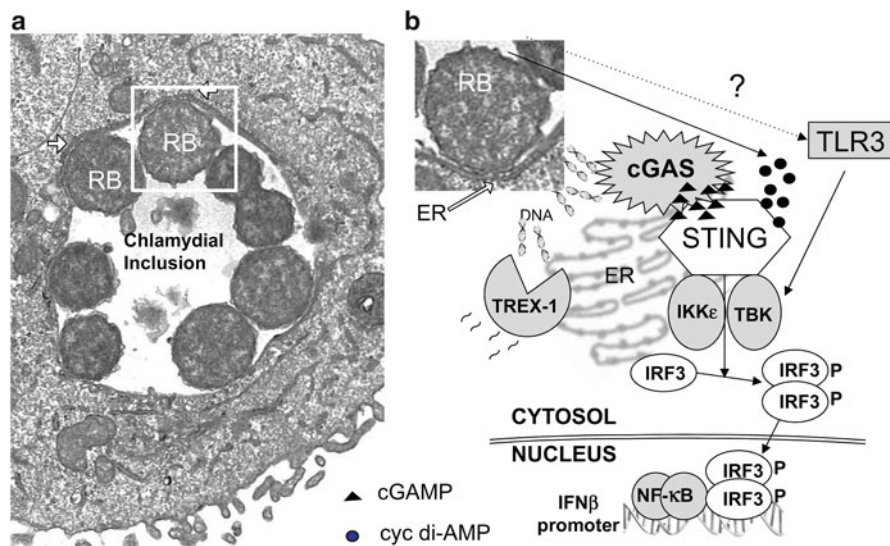
## Mechanism of IFN- $\beta$ Induction During Chlamydial Infection

Multiple host PRRs can induce IFN- $\beta$  expression during viral or bacterial infection [reviewed in [63]]. Purified *E. coli* LPS is a potent stimulator for TLR4 pathway and routinely used as a positive control for TLR4 activation [64]. However, chlamydial lipopolysaccharide (LPS) has low endotoxic activity [65, 66], which is attributed to the higher hydrophobicity of its lipid A moiety with fatty acids of longer chain length and the presence of non-hydroxylated fatty acids ester-linked to the sugar backbone. Therefore, chlamydiae stimulate TLR4 poorly, although there is one report demonstrating detection of chlamydial LPS by TLR2 [67]. Besides LPS, other bacterial ligands can stimulate TLR4, as in the case of purified hsp60 from *Chlamydia* spp. [68] However, during chlamydial infection, there is limited role for TLR4 in IFN- $\beta$  induction [41]. Cell invasion and intracellular growth is a prerequisite for IFN response during chlamydial infection. This prerequisite would suggest that intracellular receptors would be preferred over membrane-expressed receptors during infection.

Early studies showed that *C. muridarum*-induced IFN- $\beta$  is independent of TLR2 and TLR4, and some contribution from MyD88 pathway was suggested [41]. However, no contribution of TLR7 and TLR9 in IFN- $\beta$  expression was observed and TLR4-MyD88 double knockout macrophages induced similar levels of IFN- $\beta$  compared to WT macrophages [69]. Further, cytosolic RNA sensing RLR and MAVS pathways were dispensable for this response [69]. This study went on to show that the adaptor for DNA sensing, STING was essential for IFN- $\beta$  induction during *C. muridarum* infection in both mouse and human epithelial cells [69]. STING was found to localize in close proximity to the inclusion [69]. These data suggested that DNA sensors or chlamydial cyclic di-AMP could be contributing to this response. Indeed, recently it was shown that cyclic di-AMP is produced by *C. trachomatis* EBs [70]. The contribution of second messenger cyclic di-AMP in IFN- $\beta$  expression was shown by infecting HEK293T cells overexpressing STING, and transfected with IFN- $\beta$  promoter-driven luciferase reporter construct and by using fibroblasts from STING-deficient mice. Recent studies from our laboratory involved screening of multiple DNA sensors during chlamydial infection and a predominant contribution of the DNA sensor cGAS was observed in multiple cell types in response to infection using multiple *C. trachomatis* serovar [71]. The discovery of cGAS as a requirement for IFN- $\beta$  induction during chlamydial infection suggests that chlamydial DNA is available for sensing on the cytosolic side of the inclusion membrane. In support of this, cGAS was found distinctly localized on the cytosolic side of the chlamydial inclusion membrane and significant co-localization of cGAS and STING was observed after infection.

Besides DNA and cyclic dinucleotide sensing by STING, other receptors have also been shown to contribute to *Chlamydia*-induced IFN- $\beta$ . During *C. pneumoniae* infection in HUVEC cells, signaling through MAVS was found to be essential for IRF3 activation [72]. MAVS associates with TRAF3, leading to activation of IRF transcription factors and IFN- $\beta$  expression [73]. It was reported that TLR3

contributes to IFN- $\beta$  induction in mouse oviduct epithelial cells, during *C. muridarum* infection [74]. The requirement for TLR3 for IFN- $\beta$  in a bacterial infection is unique to *Chlamydia*. However, it has not been shown how TLR3 interacts with the chlamydial inclusion and the nature of chlamydial ligand engaged is unclear. Since STING was shown to be essential for IFN- $\beta$  expression during chlamydial infection in the same cell type [69], it is unclear if there is any interaction between the two pathways. Taken together, these data suggest the use of more than one host receptors for IFN- $\beta$  induction during chlamydial infection. These differences in observation could be due to: (1) differences between *C. trachomatis* vs. *C. pneumoniae* infection, (2) use of multiple pathways in cell types tested and their ability to compensate for each other, and/or (3) infection dose. The use of multiple receptors to induce the expression of the same cytokine may not be unique to *Chlamydia* spp., as multiple receptors have been suggested to play a role in IFN- $\beta$  induction during *L. monocytogenes* infection [75–78]. The signaling of type I IFN by *Chlamydia* is summarized in Fig. 1.



**Fig. 1** Model(s) for IFN beta expression during chlamydial infection. (a) An electron micrograph of chlamydial inclusion containing metabolically active RBs. (b) An enlarged image of an RB, its interaction with the host ER outside the inclusion membrane, and the proposed model(s) for IFN $\beta$  expression during infection. At least three models have been proposed for chlamydial recognition with two demonstrating the requirement for the adaptor protein STING in IFN $\beta$  induction during infection. In the first model, chlamydial EB (not shown) produce cyc di-AMP that directly interacts with STING to result in IFN- $\beta$  induction. In the second, the host DNA sensor cGAS was found to be essential for this response, with evidence for cGAMP production during infection. In support of this model, cells lacking the exonuclease TREX-1 show enhanced IFN- $\beta$  expression during infection, implicating DNA as a ligand for this response. In an alternative third model, TLR3 knock down resulted in a decrease in IFN- $\beta$  expression in a mouse oviduct cell line. The ligand engaged and its interaction with TLR3 is unknown



## Chlamydial Ligands for IFN- $\beta$ Response

The presence of enzymes essential for cyclic-di AMP synthesis and the demonstration of the presence of cyclic-di AMP in *C. trachomatis* EB [70] indicates cyclic-di AMP as a compelling ligand for IFN $\beta$  induction. Simultaneously, the significant requirement for cGAS for *Chlamydia*-induced IFN- $\beta$  indicates DNA and cGAMP as a possible ligand for IFN $\beta$  induction during chlamydial infection [71]. Although this study does not show a direct interaction of DNA with cGAS, evidence for cGAMP production was provided by demonstration of its functional transfer. HeLa cells knocked down for cGAS or STING lose their ability to induce IFN- $\beta$  upon infection, which was surprisingly rescued following their co-culture. These data suggest that cGAMP produced in cGAS competent cells during infection can function in trans by migrating to STING<sup>+</sup> cells to induce IFN- $\beta$ . These results are based on a recent study that showed that cGAMP can cross gap junctions between epithelial cells and provide cells adjacent to an infected cell intrinsic immunity independent of IFNAR signaling [79]. cGAMP binds to the same pocket in STING as cyclic di-AMP/di-GMP, but at a much lower concentration with higher affinity [80]. Indeed, the cGAS product, 2'3'cGAMP, is a much more potent ligand of STING than all other bacterial cyclic di-nucleotides described [81]. Further, human STING is responsive only to cGAMP and unresponsive to the STING ligands CMA [82] and cyclic di-AMP/cyclic di-GMP [83], unlike mouse STING which is responsive to both cyclic dinucleotides and cGAMP [84]. These studies significantly shift the importance of cGAMP over bacterial cyclic dinucleotides during *C. trachomatis* infection in human cells. However, how chlamydial DNA is transferred to cytosol remains unclear at this point. Manzanillo et al. [85] have shown that during *Mycobacterium tuberculosis* infection, phagosomal permeabilization mediated by the bacterial ESX-1 secretion system allows cytosolic recognition pathways access to DNA [85]. Numerous studies have linked IFN- $\beta$  expression to bacterial secretion systems [78, 86–88]. Small molecule inhibitors of type III secretion system (T3SS) were shown to abrogate IFN- $\beta$  expression in *C. muridarum* infected cells [89], suggesting a similar role for chlamydial T3SS in permeabilization of inclusion membrane. Previous studies [90] have shown that chlamydial reticulate bodies (RB) make direct contact with the inclusion membrane, likely through T3SS. These could be potential permeabilization points where nucleic acids could leak into cytosol and made available for host recognition. It has been shown that *Chlamydia* hijacks the host ER and several ER proteins were found localized on inclusion membrane [91, 92]. The localization of the ER protein STING [69] and cytosolic cGAS in close proximity to the inclusion membrane suggest that STING could serve as a membrane scaffold for the interactions between DNA-cGAS to take place. An alternative source of DNA detected could be host mitochondrial DNA released following damage to mitochondria in *Chlamydia*-infected cells. This argument is countered by the observation that *Chlamydia* spp. inhibit host apoptosis and no mitochondrial damage has been observed in the first 24 h of infection [93]. However, in the environment of other innate receptor recognition and production of cytokines like TNF $\alpha$ , it

is possible that mitochondrial damage occurs during *in vivo* infection and may also contribute to DNA sensor-mediated activation.

## Future Directions and Perspectives

A fascinating feature of IFN- $\beta$  inducing pathways is the resources used by the cell to detect a wide variety of pathogens to generate this important cytokine. During evolution of the immune system, viral infections likely drove this arm of innate immunity to the complex form to which it presently exists. During infection with an intracellular bacteria, the host cells responds as it would to a viral infection, detecting cytoplasmic nucleic acids and producing IFN- $\beta$ . However, IFN- $\beta$  is insufficient to eradicate bacterial infection and not protective to the host during *in vivo* infection, as in the case of chlamydial infection. In such circumstances, one can speculate that the intracellular bacteria likely exploit this antiviral pathway to its advantage. For an STI pathogen such as *C. trachomatis* that does not cause death, this would result in a prolonged infection period in the host leading to increased transmissibility. Based on studies from the mouse model, we can predict that type I IFNs likely contribute to the persistent chlamydial infection reported in humans. This could be particularly relevant during chlamydial-viral co-infection. For instance during co-infection of *C. trachomatis* with human papilloma virus, the type I IFN response resulting from the viral infection is likely to benefit *C. trachomatis* infection. Whether this results in persistent infection for either or both pathogens is not clear, although there is some evidence for *C. trachomatis* infection to be a risk factor for persistent HPV infection [94]. Over the last decade, a lot has been learnt about type I IFN induction and its role in chlamydial infection. However, the exact molecular mechanism involved in IFN- $\beta$  mediating host pathology is unclear. Further, the interaction of multiple PRRs and their cell type-specific role needs further elucidation. Over the following decade, we expect discovery of antagonists that can potentially block the pathological arm of this innate response during infection, simultaneously enhancing a protective T cell response during chlamydial infection.

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