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# TLR7, IFN $\gamma$ , and T-bet: their roles in the development of ABCs in female-biased autoimmunity

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## Abstract

The majority of autoimmune diseases have a strong gender bias, affecting mostly females. Gender-specific factors like sex-hormones, the presence or absence of a second X chromosome, and gender-specific gut microbiota may contribute to this bias. In this review we will discuss the role of the X chromosome encoded toll-like receptor 7 (TLR7) and interferon gamma (IFN $\gamma$ ) in the development of autoimmunity. We will also review recent data indicating how these factors may affect an immune response in a gender-dependent manner.

#### Keywords

gender; autoimmunity; TLR7; T-bet; IFNgamma; ABCs

About 8% of the general population suffers from one or more autoimmune disease with a strong gender bias since more than 75% of autoimmune patients are women. It is a welldocumented fact that there are differences in immune responses between males and females [1]. Overall, the female immune system is known to give stronger immune responses to infections but this characteristic may also lead to stronger responses against autoantigens [2]. Female-bias in the development of autoimmune syndromes has also been reported in various murine models of autoimmunity including NZB/W F1, NOD, and others. Differences between the immune responses of males and females have been extensively studied and, as a result, a number of explanations have been suggested. Sex hormones, the presence of a second X chromosome, microchimerism, as well as influences of various

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environmental factors have been proposed to play a role in female-biased autoimmunity [3]. Most likely, all these factors play a role in the ability of females versus males to respond to foreign and self-antigens. However, the means by which these factors affect the immune system and create gender-dependent differences has not been established. In this review we will focus on some recent findings which suggest one phenomenon that may contribute to the differences between male and female immune and autoimmune responses.

#### Differences between male and female immune responses

A number of experiments suggest that infectious diseases affect males and females differently [2, 4-6]. In general the incidence and the severity of the infections are higher in males than in females [2, 5]. Studies performed in pdriatric patients indicate that with some exceptions, the severity and prevalence of viral infections (including mumps, measles, adenovirus, coxsackievirus and respiratory syncytial virus(RSV)) is generally higher in males [4]. It has also been demonstrated that, after vaccination, females generate higher titers of measles antibodies, and antibody titers that persist for a longer time, than males [7]. Thus overall, females exhibit more robust cell-mediated and humoral immune responses than males do.

The differences between the sexes are partially due to the presence of different sex hormones. Several studies have examined how male and female sex hormones influence the immune response. It has been shown that estrogen treatment enhances the antigenic response of PBMCs and induces the expression of intracellular but not surface TLRs [8]. Estrogen treatment has been recently demonstrated to enhance the expression of UNC93B1 and IRF5 – both of these proteins play critical roles in immune responses [9, 10]. In addition several studies have demonstrated that sex hormones affect the gut microbiome which in turn may affect the immune system [11].

The presence of the second X chromosome in females may also affect immune responses in a gender-dependent manner. Although one of the two X chromosomes is inactivated in females, this inactivation is not complete and about 15% of genes encoded on the second X chromosome escape inactivation in humans [12]. The escape from inactivation leads to the overexpression of some X-linked genes in females vs males. The X chromosome encodes several immune-associated genes: CD40L, CXCR3, OGT, FOXP3, TLR7, TLR8, IL2RG, BTK, and IL9R and thus their overexpression may influence the immune response in a sex-dependent manner.

Taken together these data indicate that multiple genes can be differentially regulated in males and female which in turn may lead to the differences in immune responses to vaccinations and/or infections.

# ABCs, TLR7 and IFN $\gamma$ in autoimmunity

The contribution of sex-hormones and the X chromosome to the development of autoimmunity has been extensively studied [3]. The influence of sex hormones has been reviewed multiple times [3, 13-15] and will not be a subject of this review. Instead we will

focus on other gender-dependent differences that influence the immune system and lead to the development of autoimmunity.

In attempts to understand the origins of the female bias in autoimmunity, we and others recently identified a subset of B cells, named Age-Associated B Cells (ABCs), that are characterized by the expression of the integrin CD11c [16, 17]. As we have previously reported, ABCs accumulate in the spleens of wild type aged female but not male mice. A subset of B cells with similar characteristics appears in the spleens of autoimmune prone mice at the onset of autoimmunity. We also demonstrated that ABCs isolated from spleens of autoimmune mice are able, after stimulation, to secrete high titers of autoantibodies, predominantly of the IgG2a (IgG2c in C57BL/6 mice but termed through this review IgG2a) isotype. In the same assay other B cells, follicular, marginal zone and B1 B cells secrete relatively low levels of such autoantibodies.

Accumulation of ABCs in both aged female and autoimmune mice led us to hypothesize that these cells may contribute to the gender-bias in the development of autoimmunity. Thus it is possible that the accumulation of ABCs in females leads to a predisposition for the onset of autoimmune disease. With this in mind, we explored the mechanisms leading to the appearance of ABCs.

In our recent published work we established that the transcription factor, T-bet, is necessary and sufficient for the appearance of CD11c+ ABCs [18]. Moreover, we discovered the mechanism that leads to the expression of high levels of T-bet in B cells. Our data indicate that synergistic engagement of the B cell antigen receptor (BCR), TLR7, and IFN $\gamma$  receptor is required for the expression of high levels of T-bet in B cells. In turn, T-bet drives expression of the ABC phenotype and a switch of antibody production to the IgG2a isotype (Figure 1) [18]. In support of this hypothesis, we also showed that intact TLR7 and IFN $\gamma$ R signaling is required for the accumulation of ABCs in aged female mice.

All of these three factors (BCR, TLR7 and IFN $\gamma$ ) have been shown to play a crucial role in the development of autoimmunity and will be further discussed in this review.

#### Role of TLR7 in autoimmunity

TLR7 is an intracellular toll-like receptor expressed by various cell types including B cells. TLR7 recognizes single-stranded RNA. Involvement of TLR7 in the development of autoimmunity has been demonstrated by multiple groups using various approaches. For example it has been reported that TLR7 deficiency leads to the ablation of autoimmunity in the MRL<sup>lpr</sup> mouse model of lupus [19]. In addition, treatment of MRL<sup>lpr</sup> mice with oligodeoxynucleotides with immunoregulatory sequences (IRS), that specifically block signaling via TLR7, has been reported to attenuate the disease [20]. Our group has recently demonstrated that TLR7 deficiency also abolishes the appearance of autoantibodies in B6.Nba2 and MER<sup>-/-</sup> murine models of SLE [21].

Several studies reported that the number of copies of the Tlr7 gene affects development of autoimmunity. Thus, duplication of Tlr7 in autoimmune-prone mice has been shown to accelerate the development of lupus [22] and overexpression of Tlr7 in non-autoimmune

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mouse strains leads to the development of spontaneous autoimmunity [23]. Moreover, Hwang et al., have recently demonstrated that overexpression of *Tlr7* by B cells specifically, is sufficient to drive production of autoantibodies, indicating a direct effect of TLR7 signaling in B cells [24]. Looking more closely at the mechanism behind TLR7-induced autoimmunity, Walsh et al. reported that overexpression of *Tlr7* leads to the development of spontaneous germinal centers and promotes differentiation of plasmablasts in a T cell dependent manner [25].

In addition to animal models, TLR7 is also involved in the development of autoimmunity in human patients. After it was demonstrated that TLRs are involved in the pathogeneses of murine lupus like disease, several studies examined the levels of the expression of these molecules in leukocytes in human SLE. A study performed on SLE patients by Komatsuda et al., demonstrated elevated expression levels of TLR7 in SLE patients [26]. Additionally, a significant increase in *TLR7* gene copy number was detected in SLE patients compared to healthy controls [27]. However, Kelley et al. could not replicate this result and reported no significant difference in *TLR7* copy number between SLE patients and healthy controls [28].

Together these data implicate TLR7 as a potential regulator of autoimmunity. Moreover, the copy number of the Tlr7 gene may be an important contributory factor. However, does TLR7 affect the immune response in a gender-dependent manner? Localization of Tlr7 gene to the X chromosome in both mice and humans implies that its expression might be regulated in gender-dependent manner. In mammals, females have two X chromosomes while males have only one X and one Y chromosome. In order to have equal levels of gene expression from X chromosomes in both genders, one X chromosome in female cells is inactivated. However, some genes on the X chromosome escape inactivation leading to potentially higher expression levels of several X-linked genes in females. In fact, the number of genes escaping inactivation is quite high, reaching ~15% of X-linked genes in humans [12]. Thus, localization of Tlr7 gene to the X chromosome may potentially lead to its overexpression in females. The importance of X-linked gene dosage for the development of SLE in humans has been proposed based on the fact that males with Klinefelter's syndrome (males with an XXY genotype) have a greater risk for the development of lupus compared to the general male population [29]. In fact, these males are as susceptible to SLE as the general female population, indicating that despite differences in hormonal content, the presence of a second X chromosome is crucial for female-biased development of SLE.

In summary, the data acquired by multiple groups in both murine and human autoimmunity indicate that TLR7 is critical for the development of disease. Moreover, the level of TLR7 expression plays an important role and might differ in a gender-dependent manner due to the location of the *Tlr7* gene on the X chromosome. However, none of these studies have provided a mechanism to explain how TLR7 signaling drives the production of pathogenic autoantibodies.

### Interferon gamma and autoimmunity

The effect of IFN $\gamma$  on the development of autoimmunity is well established. It has been demonstrated that IFN $\gamma$  is essential for the development of autoimmunity in MRL<sup>lpr</sup> mice

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[30-32]. Deletion of IFN $\gamma$  abolishes the appearance of autoantibodies in NZBxW F1 mice [33] and treatment of NZBxW F1 mice with blocking anti-IFN $\gamma$  antibodies has been demonstrated to ameliorate the disease [34]. In addition it has been demonstrated by Seery et al., that IFN $\gamma$  transgenic mice, mice that overexpress IFN $\gamma$ , develop a lupus-like syndrome [35]. The authors demonstrated the production of high levels of anti-nucleosome, antihistone, and anti-dsDNA antibodies by female transgenic mice. It is worth noticing that male transgenic mice did not develop a lupus-like phenotype [35]. Another piece of evidence came from *Roquin*<sup>san/san</sup> mice in which a mutation of ROQUIN leads to reduced decay of IFN $\gamma$  mRNA resulting in increased numbers of Tfh cells, germinal center B cells, and autoantibodies [36]. Notably, IFN $\gamma$ R deficiency prevents autoimmunity in Roquin<sup>san/san</sup> mice, confirming a crucial role of IFN $\gamma$  in this model of SLE.

With regards to human autoimmunity, increased levels of IFN $\gamma$  have been reported for patients with SLE, systemic sclerosis and Sjorgen's syndrome [37-40]. Peripheral blood monocytes (PBMCs) from patients with multiple sclerosis (MS) were demonstrated to produce more IFN $\gamma$  upon in vitro stimulation with a peptide from proteolipid protein (PLP) when compared to healthy controls [41]. Moreover, the level of IFN $\gamma$  production was gender-dependent. Female cells produced higher levels of IFN $\gamma$  than male cells in both MS and healthy control groups. Gender-related differences in IFN $\gamma$  production has also been reported in other systems. For example, female mice have been demonstrated to produce higher levels of IFN $\gamma$  in response to *Paracoccidioides brasiliensis* infection when compared to males [42]. In this case sex hormones were shown to be responsible for the effect. Sexdetermined susceptibility to *Leishmania Mexicana* has also been explained by higher levels of IFN $\gamma$  production by females, compared to males [43].

Taken together these data indicate that IFN $\gamma$  plays a critical role during the development of autoimmunity in both murine models and in human patients. Since females have been reported to produce higher levels of IFN $\gamma$  under a number of circumstances, it is worth suggesting that IFN $\gamma$  might be one of the factors contributing to the female bias in autoimmunity.

#### Role of T-bet in autoimmunity

The data we have reviewed above clearly point to a critical role for TLR7 and IFN $\gamma$  in the development of autoimmunity. Moreover, gender related differences might apply to the expression of TLR7 and IFN $\gamma$  production, suggesting that these factors might be involved in the female-bias of autoimmunity. However, none of the above studies have explained how IFN $\gamma$  or TLR7 deficiency affects autoimmune disease. Since our data indicate that signaling via IFN $\gamma$ R and TLR7 are required for T-bet induction in B cells (Figure 1), we hypothesized that ablation of ABCs by TLR7 or IFN $\gamma$  deficiency might lead to amelioration of autoimmunity. We have demonstrated that, at least in MER<sup>-/-</sup> mice, deletion of TLR7 prevents the accumulation of T-bet<sup>+</sup> ABCs and this phenomenon correlates with the absence of autoantibodies [21]. However this hypothesis has to be further tested directly by depletion of T-bet<sup>+</sup> B cells in autoimmune mice. The progression of autoimmunity in mice with B cell specific deletion of T-bet has never been studied. This is a very important experiment which

we will perform in the near future. However, the role of T-bet expression during the progression of autoimmunity will be briefly reviewed here.

T-bet deficiency leads to different outcomes depending on the autoimmune model in question. T-bet deletion ameliorates disease in some models of autoimmunity including collagen antibody induced arthritis (CAIA) [44],myelin oligodendrocyte glycoprotein (MOG) induced experimental autoimmune encephalomyelitis (EAE) [45], Type 1 Diabetes (T1D) in NOD mice [46], and lupus like disease in MRL<sup>lpr</sup> animals [47]. In contrast, worsening of proteoglycal or *S. aureus* induced arthritis has been reported in T-bet–/– mice [48]. All these studies are complicated by the fact that the diseases in question were studied in mice in which all cell types lacked T-bet expression. Since T-bet is involved in the functions of many different cell types (T cells, NK cells, DCs, B cells), it is difficult to know which cell types are responsible for the improvement or worsening of the disease in the absence of T-bet. For this reason we believe that cell type-specific deletion of T-bet will shed more light on its role in autoimmunity.

## **Concluding remarks**

The data reviewed above indicate that TLR7 and IFN $\gamma$  might be amongst the key factors that drive gender-related differences in immune and autoimmune responses. More studies are required in order to clarify the role of T-bet expressing B cells in gender-biased autoimmunity. For instance, the role of T-bet expression in B cells during the onset of autoimmunity has to be determined. In addition, it has to be revealed whether males and females differ in their abilities to induce T-bet expression in B cells and even whether TLR7 is expressed at different levels in the lymphocytes of males and females. We think that answers to these questions might lead us to a better understanding of the mechanisms of female bias in autoimmunity and might lead ultimately to the development of better and perhaps gender specific therapies and/or diagnostics for these illnesses.

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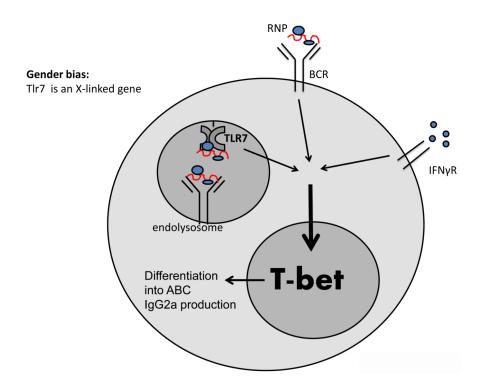
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#### **HIGHLIGHTS**

- TLR7, IFN $\gamma$  and BCR synergistically induce expression of T-bet in B cells

- T-bet expressing B cells acquire distinct phenotype and become ABCs
- accumulation of ABCs in females may lead to a predisposition for the autoimmunity
- TLR7 and IFNy may drive gender-related differences in (auto)immune response



#### Figure 1. Model for T-bet induction in B cells

Synergistic signaling via B cell antigen receptor (BCR), Toll-like receptor 7 (TLR7), and IFN $\gamma$ R in B cells leads to the induction of high levels of T-bet expression, which in turn drive the expression of an ABC phenotype and class switching to the production of IgG2a antibodies. A possible explanation for the gender bias of autoimmunity: *TLR7* is an X-linked gene, which might be overexpressed in females compared to males due to incomplete X-inactivation. Overexpression of *TLR7* and/or accumulation of self-antigen (RNPs) due to the ineffective clearance of apoptotic cells may lead to the accumulation of ABCs and the development of autoimmunity.