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## ABSTRACT

In Sjögren's Syndrome (SS), inherent glandular defects, autoimmunity, and mononuclear cell infiltration within the salivary glands cause reduced salivation leading to xerostomia. Excessive production of type I interferons (IFN), triggered by environmental and genetic factors, is considered pathogenic in this disorder. However, whether type I IFN production is causative or an outcome of the disease process is not known. To address this question, we introduced a deficiency of interferon alpha receptor 1 (*Ifnar1*) into B6.*Aec1Aec2* mice, which are known to have the genetic loci necessary for developing a SS-like disorder. This new mouse strain, B6.*Aec1Aec2Ifnar1*<sup>-/-</sup>, lacking type I IFN-mediated signaling, was characterized for pilocarpine-induced salivation, the presence of serum autoantibodies, sialoadenitis, and dacryoadenitis. Compared with the B6.*Aec1Aec2Ifnar1*<sup>+/+</sup> (wild-type) mice, the B6.*Aec1Aec2Ifnar1*<sup>-/-</sup> (knockout) mice had significantly lower mononuclear cell infiltration in the salivary and lacrimal glands. The knockout mice were completely protected from salivary gland dysfunction. Surprisingly, they had a robust autoantibody response comparable with that of the wild-type mice. These findings demonstrate that, in the absence of type I IFN-mediated signaling, systemic autoantibody responses can be dissociated from glandular pathology. Our study suggests that, in genetically susceptible individuals, the type I IFN pathway can instigate certain features of SS.

**KEY WORDS:** xerostomia, sialoadenitis, autoimmunity, salivary gland, mouse, interferons.

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# Type I Interferon Receptor Deficiency Prevents Murine Sjögren's Syndrome

## INTRODUCTION

Xerostomia or dry mouth is one of the characteristic features of Sjögren's Syndrome (SS) and affects several aspects of oral health (Carr *et al.*, 2012). The lack of sufficient saliva often leads to an increased incidence of dental caries and periodontal disease, necessitating frequent interventions by a dental professional. Moreover, dry mouth causes difficulty in speaking and the chewing and swallowing of food and leads to a decreased quality of life.

Salivary gland dysfunction in SS has been attributed to a systemic as well as a localized autoimmune response within the salivary glands (Kang *et al.*, 2011). While the systemic autoimmunity in SS is highlighted by the detection of autoantibodies in sera, the presence of inflammatory cells within the salivary glands is indicative of a localized immune response in the disease. Although it has been difficult to pinpoint the precise triggers responsible for initiating SS, viral infections causing excessive production of type I interferons (IFNs) have long been the suspects (Vakaloglou and Mavragani, 2011). Type I IFNs consist of several IFN proteins that bind and signal through a common interferon alpha receptor (IFNAR) (Trinchieri, 2010). Engagement of IFNAR eventually leads to the activation of a multitude of genes, known as the IFN-responsive genes, which affect several aspects of cellular metabolism. Increased expression levels of IFN-responsive genes in SS patients have led to the suggestion that type I IFNs play an important role in the pathogenesis of this disease (Hjelmervik *et al.*, 2005; Emamian *et al.*, 2009). Paradoxically, reduced levels of circulating IFN- $\alpha$  in SS patients have been used as a rationale for clinical trials with IFN- $\alpha$ . However, these trials have reported variable effects on the disease (Shirota *et al.*, 2008). Thus, whether type I IFN response is causative for SS is difficult to determine in patients.

Previous work from our laboratory has demonstrated that activation of the type I IFN pathway through the Toll-like receptor 3 (TLR3) agonist poly(I:C) rapidly leads to reversible salivary gland dysfunction in mice (Deshmukh *et al.*, 2009). This dysfunction did not occur in mice lacking IFNAR (Nandula *et al.*, 2013). In addition, poly(I:C) treatment accelerated and increased the severity of sialoadenitis in mice genetically susceptible to the development of SS-like disease (Nandula *et al.*, 2011). While these studies demonstrate the role played by TLR3 activation in the induction of salivary gland dysfunction and disease, we were interested in investigating whether abrogation of type I IFN signaling can prevent the development of SS-like disease. To address this question, in this study, we developed a novel genetic mouse model that lacks the IFNAR (thereby abrogating all signaling through type I IFNs), but that has all the genes necessary for the development of SS-like disorder.

Currently, there are several mouse models for SS described in the literature, and each model has its advantages and disadvantages for investigating this complicated disorder (Delaleu *et al.*, 2011). Since the IFNAR deficiency was on the C57BL/6 (B6) background (Yasuda *et al.*, 2007), we used the B6.*Aec1Aec2* mouse for our study. The B6.*Aec1Aec2* spontaneous mouse model for SS-like disorder recapitulates certain features of the human disease (Cha *et al.*, 2002). It carries 2 genetic regions, *Idd5* and *Idd3*, from the non-obese diabetic (NOD) mouse. The advantages of using this model are that it is spontaneous and characterized by the presence of circulating autoantibodies, lymphocytic foci within the salivary glands, and a progressive loss of glandular function. However, unlike the human disorder, there is no female-to-male preponderance of disease. In this study, we generated a novel strain, the B6.*Aec1Aec2Ifnar1*<sup>-/-</sup> knockout (KO) mouse, and investigated the development of SS-like disorder in this mouse.

## MATERIALS & METHODS

### Animals

All mice used in this study were bred and maintained under specific pathogen-free (SPF) conditions in the University of Virginia vivarium. All experiments were approved by the Institutional Animal Care and Use Committee, and all protocols followed National Institutes of Health guidelines for humane practices. The B6.*Aec1Aec2* mice (Cha *et al.*, 2002) were crossed with the B6.*Ifnar1*<sup>-/-</sup> mice (Yasuda *et al.*, 2007) to generate F1 mice. The F1 mice were intercrossed and backcrossed to parental strains to generate the B6.*Aec1Aec2Ifnar1*<sup>-/-</sup> mouse. All progeny were genotyped for the *Aec1* and *Aec2* loci using a panel of microsatellite markers as described previously (Nguyen *et al.*, 2006). The *Ifnar1* deficiency was monitored by a gene-specific polymerase chain-reaction (PCR) and flow cytometry. Pilocarpine-induced salivation was measured as described previously (Deshmukh *et al.*, 2009). All experiments were performed in female mice.

### Gene Expression Analysis

Expression of IFN-responsive genes in submandibular glands was determined by quantitative PCR with Taqman probes as described previously (Deshmukh *et al.*, 2009). A pooled RNA sample from submandibular glands of age-matched B6 mice was used as a calibrator to calculate relative gene expression by the 2<sup>-ΔΔCt</sup> method (Livak and Schmittgen, 2001).

### Antibody Analysis

Immunoglobulin levels in sera were measured by ELISA as described earlier (Bagavant *et al.*, 2002). Serum autoantibodies to a mouse submandibular gland cell line SCA-9-15 (ATCC, Manassas, VA, USA) extract were analyzed by Western blotting as described previously, with few modifications (Deshmukh *et al.*, 1999). Goat anti-mouse IgG coupled to IRDye800 (LI-COR Biotechnology, Lincoln, NE, USA) was used as secondary antibody. Bound antibody was visualized with the ODYSSEY CLx infrared imaging system following manufacturer's directions,

and intensity of staining was quantitated with Image Studio 2.1 software (LI-COR Biotechnology). Anti-nuclear antibodies (ANA) in serum were detected by indirect immunofluorescence assay with HeLa cells as previously described (Deshmukh *et al.*, 1999).

### Stereology

Submandibular salivary gland and lacrimal gland pieces were fixed in 10% buffered formalin and embedded in paraffin, and sections were stained by hematoxylin and eosin. Quantitative analyses of glandular inflammation were done with Stereoinvestigator software v3.0 (MBF Bioscience, Williston, VT, USA) (Glaser and Glaser, 2000). Briefly, a contour was traced around each piece of gland, and a sampling grid of 400 μm x 400 μm was randomly placed over the contour. Using the area fraction fractionator probe, we obtained a systematic random sample by positioning a counting frame of 300 μm x 250 μm containing markers at 10-micron intervals. Each marker was interrogated for the presence or absence of inflammatory cells, and the total gland area and inflammatory focus area were estimated. Salivary gland areas of 10.4 ± 0.5 x 10<sup>6</sup> sq microns and lacrimal gland areas of 3.3 ± 0.4 x 10<sup>6</sup> sq microns were studied for each mouse. Disease scores were calculated as (inflammatory area/total gland area) x100 for each mouse.

### Flow Cytometry

Analysis of IFNAR expression on peripheral blood cells was carried out by flow cytometry according to standard protocols. Experimental details are provided in the Appendix.

## RESULTS

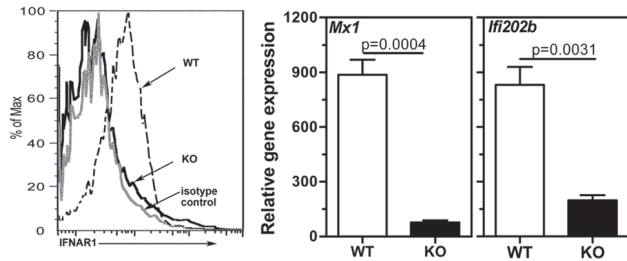
### Expression of IFN-responsive Genes is Suppressed in B6.*Aec1Aec2Ifnar1*<sup>-/-</sup> Mice

The *Ifnar1* mutation in C57BL/6 mice was moved to the B6.*Aec1Aec2Ifnar1*<sup>+/+</sup> mice (WT) to generate B6.*Aec1Aec2Ifnar1*<sup>-/-</sup> mice (KO). Appendix Fig. 1 shows the PCR products used to characterize the mouse genotype. To confirm the phenotype, we analyzed expression of IFNAR on peripheral blood mononuclear cells by flow cytometry (Fig. 1, left panel). IFNAR expression was not detected on cells obtained from the KO mice.

The WT mice showed spontaneous up-regulation of type I IFN-responsive genes in their salivary glands (Nguyen *et al.*, 2009). Therefore, expression of the IFN-responsive genes *Mx1* and *Ifi202b* was analyzed in submandibular glands of WT and KO mice at 8 wks of age. The expression of both genes was significantly suppressed in the KO mice (Fig. 1, right panel).

### KO Mice are Protected from Salivary Gland Dysfunction

Spontaneous loss of salivary gland function is one of the major characteristics of the SS-like disorder in the B6.*Aec1Aec2* mouse. To determine the effect of IFNAR deficiency on this phenotype, we measured pilocarpine-induced saliva at different time-points in the WT and KO mice (Fig. 2). At an early age



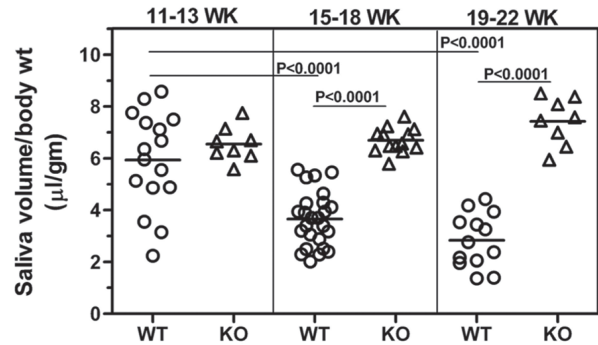
**Figure 1.** Lack of IFNAR in *B6Aec1Aec2* mice suppresses the expression of type I IFN-responsive genes *Mx1* and *Ifi202b*. **(Left panel)** Flow cytometry analysis showing a representative histogram of IFNAR staining on PBMCs obtained from WT (dotted line) and KO (solid line) mice. The staining by isotype control is shown by a gray line. **(Right panel)** Gene expression levels of *Mx1* and *Ifi202b* in the submandibular glands obtained from WT (n = 4) and KO (n = 3) mice at 8 wks of age were determined by real-time PCR with Taqman primers. The data are represented as mean  $\pm$  SEM relative gene expression over a pooled RNA sample from an age-matched C57BL/6 mouse, which was used as the calibrator. *Gapdh* was used to normalize the expression. Statistical significance was determined by the Mann-Whitney test. Similar results were obtained in an additional experiment.

(11-13 wks), the difference in mean saliva volume between the WT and KO mice was not statistically significant. At later ages, there was a progressive drop in saliva volumes in the WT mice. In contrast, the mean saliva volumes in the KO mice, monitored up to 22 wks, did not decrease. At 15 to 18 wks and 19 to 22 wks, differences in mean saliva volumes between WT and KO mice were highly significant ( $p < .0001$ ). Analysis of these data clearly demonstrates that genetic ablation of type I IFN signaling in the *B6.Aec1Aec2* mice protects them from salivary gland dysfunction, as indexed by a ratio of salivary volume to body weight.

### IFNAR Deficiency Does Not Affect Systemic Autoimmune Response

The IFNAR deficiency did not dramatically alter the total serum IgG and IgM levels (Appendix Fig. 2). Although in the WT mice at 7 to 10 wks of age, the mean IgM levels were significantly higher ( $p = .0101$ ), at later time-points the differences between the 2 groups were not statistically significant. The mean IgG levels between WT and KO mice, at both early and late ages, were not significantly different.

To determine whether the protection from salivary gland dysfunction in KO mice was tied to suppression of spontaneous autoimmune responses, we analyzed the presence of ANA in sera obtained from WT and KO mice at different time-points. There was no difference in the incidence or intensity of ANA staining between the WT and KO mice at all time-points tested (Fig. 3, left panel, and Appendix Fig. 3, showing a representative ANA staining). A similar trend was observed when sera were tested for reactivity to a mouse submandibular gland cell line SCA-9-15 (data not shown). Serum reactivity to salivary gland antigens was also analyzed in Western blots (Fig. 3, right panel). Autoantibodies reactive with multiple proteins were detected in the sera of both WT and KO mice, and the intensities of staining were not different between the groups. Analysis of



**Figure 2.** Saliva production decreases over time in WT mice but not in KO mice that lack the type I IFN signaling pathway. Pilocarpine-induced saliva production was measured at the different ages indicated in the Fig. Data are represented as the ratio of saliva volume ( $\mu$ l) to body weight (gm). Student's *t* test was used to determine statistical significance. Saliva volumes (mean  $\pm$  SEM) in WT mice at 15 to 18 wks ( $3.65 \pm 0.21$ ) and 19 to 22 wks ( $2.83 \pm 0.28$ ) were significantly lower compared with those at 11 to 13 wks ( $5.93 \pm 0.46$ ) of age, indicating a progressive loss of function. In contrast, in KO mice, the mean saliva volumes between early and late ages were not different. Although WT and KO mice had comparable saliva production at 11 to 13 wks of age, the differences between the 2 groups were highly significant at the 15 to 18-week ( $p < .0001$ ) and 19- to 22-week ( $p < .0001$ ) time-points.

these data indicates that IFNAR deficiency in the *B6Aec1Aec2* mice did not permanently affect total immunoglobulin levels nor did it affect autoantibody reactivity to multiple cellular antigens. Overall, these findings are consistent with the data showing no differences in T-, B-, and NK cell populations between WT and KO mice (Appendix Fig. 4).

### IFNAR Deficiency Affects Inflammatory Cell Infiltration

H&E-stained sections of submandibular glands obtained from WT and KO mice were studied for the presence of inflammatory cell infiltrates at 20 to 22 wks, and representative histopathology is shown in Figs. 4A-4D. Quantitative analyses showed larger areas of focal inflammation in WT mice ( $8.49 \pm 1.7 \times 10^4$  sq microns) compared with KO mice ( $3.8 \pm 1.8 \times 10^4$  sq microns). The inflammatory area fraction of the total glandular tissue studied was calculated for each mouse and is represented as severity of sialoadenitis (Fig. 4, lower panel). The mean inflammatory area fraction was significantly lower ( $p = .0022$ ) in the KO mice ( $0.37 \pm 0.2$ ) compared with that in age-matched WT mice ( $0.75 \pm 0.13$ ). Lacrimal gland inflammation (Figs. 4E-4H) was also reduced in KO mice ( $1.7 \pm 1.2 \times 10^4$  sq microns) compared with that in WT mice ( $10.0 \pm 2.9 \times 10^4$  sq microns). Similarly, the lacrimal inflammatory area fraction was significantly lower ( $p = .0003$ ) in the KO mice ( $0.56 \pm 0.4$ ) compared with that in age-matched WT mice ( $3.43 \pm 1.08$ ).

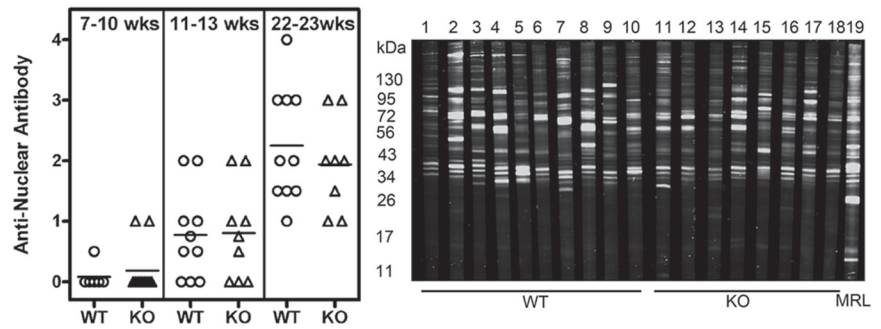
## DISCUSSION

Type I IFNs are pluripotent cytokines that play a crucial role in host defense against viral infections (Isaacs and Lindenmann,

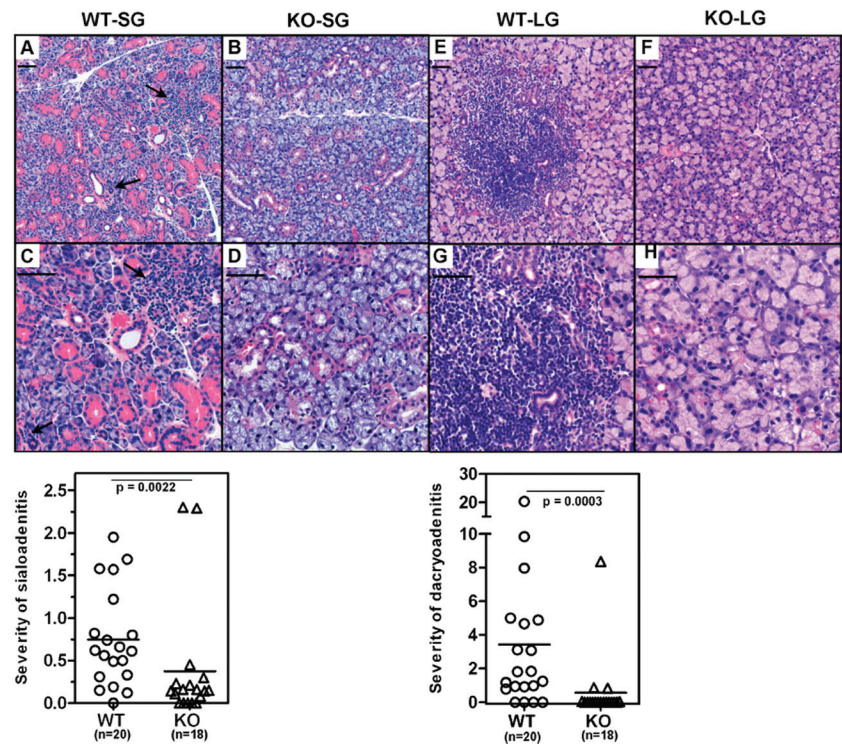


1957). However, they have also been implicated in the pathogenesis of autoimmune disorders such as lupus and SS (Trinchieri, 2010). Although the up-regulation of IFN-responsive genes in SS patients suggests a possible role for type I IFNs in the pathogenesis of this disease, whether they are involved in the initiation or perpetuation of SS is not known. This study, in a spontaneous mouse model for SS-like disorder, clearly demonstrates that lack of type I IFN signaling prevents the development of disease. Mice deficient for *Ifnar1*, but harboring other genes necessary for the development of SS-like disorder, were protected from both mononuclear cell infiltration within the lacrimal and salivary glands and the loss of saliva production. Interestingly, the KO mice had mounted a robust systemic autoimmune response, thereby dissociating the systemic humoral autoimmune response from salivary gland dysfunction.

In our model, the mice were protected from salivary gland dysfunction, yet they had a robust systemic autoimmune response. This suggests that systemic autoimmune responses need not be under the control of the type I IFN pathway. The effect of IFNAR deficiency has been studied in other models of systemic autoimmunity, particularly in lupus (Santiago-Raber *et al.*, 2003; Agrawal *et al.*, 2009). Although some lupus mouse models showed that the lack of *Ifnar1* was responsible for decreasing autoantibody levels, others, like the *FcγRIIB<sup>-/-</sup>Yaa Ifnar1<sup>-/-</sup>* mice, showed that type I IFN signaling did not influence the levels of ANA and anti-Sm autoantibodies (Richez *et al.*, 2010). Similar to our finding, the total immunoglobulin levels were not perturbed in these mice. These observations have a considerable bearing on our understanding of the pathogenic mechanisms operative in SS. The type I IFN pathway has been implicated for initiating and amplifying a systemic autoimmune response, by exerting effects on antigen-presenting cells and by influencing B-cell activation (Voulgarelis and Tzioufas, 2010). However, while depletion of B-cells with Rituximab therapy increased the expression of type I IFN-responsive genes in rheumatoid arthritis patients, in SS patients, it had no effect on the IFN signature (Vosslander *et al.*,



**Figure 3.** Both WT and KO mice generated a robust autoantibody response. Sera obtained from WT and KO mice at different time-points were analyzed for the presence of autoantibodies. (Left panel) Intensity of ANA staining by indirect immunofluorescence. Representative ANA pictures are shown in Appendix Fig. 3. The incidence and severity of ANA staining were not different between the WT and KO mice. (Right panel) A representative image of a Western blot showing reactivity to proteins from a mouse submandibular gland cell line. All sera were used at 1:200 dilutions, and goat anti-mouse IgG antibody coupled to IRDye 800CW was used to detect bound antibodies. Lanes 1 to 10 are sera from individual WT mice, lanes 11 to 18 are sera from individual KO mice, and lane 19 has pooled sera from lupus-prone MRL *lpr/lpr* mice. Similar results were obtained in an additional experiment.



**Figure 4.** The IFNAR deficiency suppresses mononuclear cell infiltration within the submandibular and lacrimal glands. Submandibular (A-D) and lacrimal glands (E-H) obtained from WT (n = 20) and KO (n = 18) mice were evaluated for mononuclear cell infiltration in H&E-stained sections. The inflammation was quantitated by an individual blinded to experimental details. A, B, E, and F show representative low-magnification images, and C, D, G, and H are images at higher magnification. Severity of sialoadenitis and dacryoadenitis (lower panels) represents the fraction of area covered by inflammatory infiltrates in salivary and lacrimal glands, respectively, for each mouse. Statistical significance was calculated by the Mann-Whitney test. The mean severity scores of sialoadenitis and dacryoadenitis in the WT mice were significantly higher than those in the KO mice. The arrows indicate the foci of mononuclear cell infiltration. Scale bar = 50 microns.

2011; St. Clair *et al.*, 2013). Considering these observations together with our results from the KO mice, we suggest that the influence of the type I IFN pathway on autoantibody production might be dependent on the disease type and interaction with other genetic factors.

In B6.*Aec1Aec2* mice, IFNAR deficiency prevents inflammatory cell infiltration in the salivary glands. Several mechanisms might be operative in exerting this protective effect. Type I IFNs are known to influence the production of chemokines (Lang *et al.*, 2006). Our previous work has shown that activation of the type I IFN pathway leads to up-regulated gene expression for several chemokines within the submandibular glands (Nandula *et al.*, 2011). This causes accelerated mononuclear cell infiltration within the salivary glands. A recent study suggests that 2 chemokines, CXCL10 and CXCL12, expressed in the salivary glands of B6.*Aec1Aec2* mice and SS patients are responsible for the progression of disease (Horvath *et al.*, 2012). Thus, lack of type I IFN signaling in the KO mice might decrease the localized production of such chemokines and thereby reduce inflammatory cell infiltration within the salivary glands. Alternatively or in addition, the expression patterns of different chemokine receptors on inflammatory cells can be altered in the absence of type I IFN signaling, thereby affecting their migratory properties. A systematic comparison of changes in chemokine and chemokine receptor expression between the KO and WT mice will yield targets for interfering with disease progression in SS. Indeed, administration of a CXCL10 antagonist to MRL *lpr/lpr* mice resulted in decreased salivary gland inflammation (Hasegawa *et al.*, 2006).

In summary, our study demonstrates that type I IFN is responsible for initiating features of SS-like disorder in B6.*Aec1Aec2* mice. This study supports the thesis that, in genetically susceptible individuals, up-regulation of type I IFN production can lead to the development of SS.

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