

FIGURE 2. Infiltrating cells in salivary glands isolated from RORγt Tg mice. **(A)** Flow cytometric analysis of lymphocyte-gated cells (15–20%) stained with CD19, CD3, CD4, and CD8 on CD3⁺ subsets in infiltrated cells isolated from the salivary glands of RORγt Tg mice. Four mice per group were analyzed and representative data are shown. **(B)** Immunofluorescence analysis of salivary glands from RORγt Tg mice at 4, 6, and 12 wk of age. Data are representative of two tissue samples with similar results. **(C)** Apoptotic cells in salivary glands of C57BL/6 and RORγt Tg mice detected by the TUNEL method. Data are representative of three tissue samples with similar results. Original magnification ×100 (B) and ×600 (C).

glands of RORγt Tg mice compared with C57BL/6 mice (Fig. 4A). Second, overexpression of Th1 (*Ifng*, *Il2*, *Il12*, *Tnf*, *Tbet*)-, Th2 (*Il4*, *Gata3*)-, and Tfh (*Il21*, *Bcl6*)-related factors was also noted in the inflammatory lesions of salivary glands of RORγt Tg mice compared with C57BL/6 mice (Fig. 4B). Comparison of the expression of

transcriptional factors related to Th cells in CD4⁺ T cells of salivary glands of RORγt Tg mice showed significantly higher expression of *Rorc*, *Tbet*, *Gata3*, and *Bcl6* in CD4⁺ T cells than those of the spleen (Fig. 4C). Moreover, infiltrating CD4⁺ T cells isolated from salivary glands of RORγt Tg mice produced IL-17, INF-γ, IL-4, and IL-21

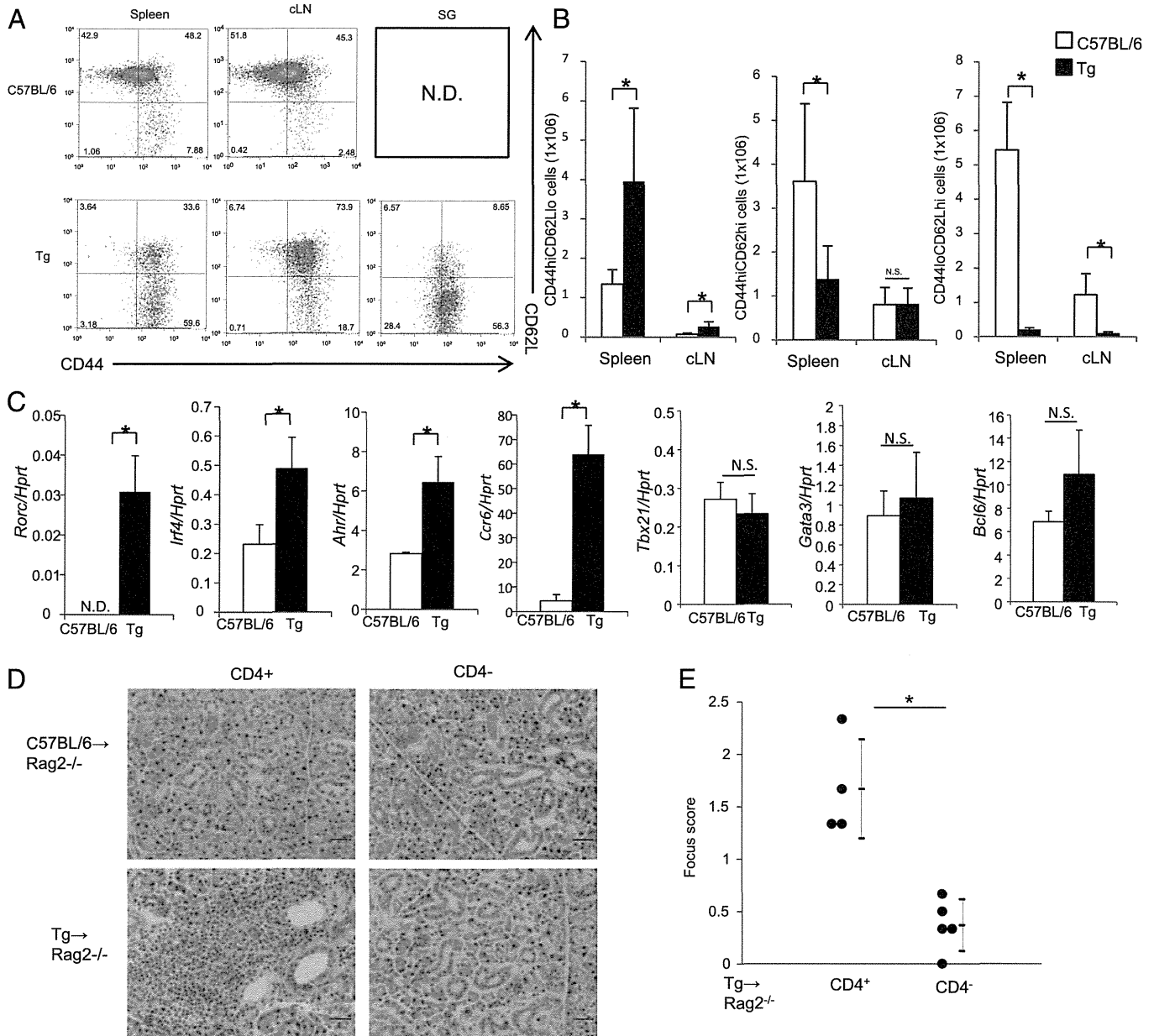


FIGURE 3. ROR γ T-overexpressed CD4⁺ T cells in ROR γ T Tg mice. **(A)** Flow cytometric analysis of naive and effector CD4⁺ T cells in spleen, cLN, and salivary glands of C57BL/6 and ROR γ T Tg mice. Five mice per group were analyzed and representative data are shown. **(B)** Mean number of CD44^{hi}CD62L^{lo}, CD44^{hi}CD62L^{hi}, and CD44^{lo}CD62L^{hi} cells gated on CD4⁺ cells from spleen and cLN of C57BL/6 and ROR γ T Tg mice. Four mice per group were analyzed and representative data of two independent experiments with consistent results are shown. **(C)** Quantitative PCR analysis of mRNA expression in splenic CD4⁺ T cells isolated from C57BL/6 and ROR γ T Tg mice. The experiment was performed in duplicate. Data were normalized to the expression of the reference gene, *Hprt*. **(D)** Purified splenic CD4⁺ T and CD4⁻ cells (1×10^7 cells) were inoculated into Rag2^{-/-} mice. At 22 wk postinoculation, the pathology of salivary glands was analyzed. Salivary glands were sectioned at 4 μ m, and each section was stained with H&E. Scale bars, 30 μ m. Data are representative of four tissue samples with similar results. **(E)** Histological score of inflammatory lesions in salivary glands of ROR γ T Tg CD4⁺ or CD4⁻ → Rag2^{-/-} mice at 22 wk postinoculation. * $p < 0.05$ (Mann–Whitney *U* test). N.D., not detected.

(Fig. 4D). These data suggest the involvement of not only Th17 but also Th1, Th2, and Tfh cells in the pathogenesis of sialadenitis.

IL-17 is not essential for the induction of sialadenitis in ROR γ T Tg mice

To determine the significance of IL-17 in the development of sialadenitis, we generated IL-17-deficient ROR γ T Tg mice (IL-17^{-/-}/Tg). Sialadenitis occurred spontaneously in IL-17^{-/-}/Tg mice and was similar to that in ROR γ T Tg mice (Fig. 5A). Furthermore, the focus score of inflammatory lesions in the salivary glands (Fig. 5B) and the saliva flow (Fig. 5C) of these mice was comparable to that of ROR γ T Tg mice. Immunoflu-

orescence (Fig. 5D) and flow cytometric analyses (Fig. 5E) of infiltrated cells in salivary glands showed the presence of CD4⁺ T and B cells in the salivary glands of IL-17^{-/-}/Tg mice. The percentages of infiltrating CD3, CD19, CD4, and CD8 T cells in IL-17^{-/-}/Tg mice were not different from those counted in ROR γ T Tg mice (Fig. 5E). Quantitative PCR showed that mRNA expression of *Il17a* was absent in salivary glands of IL-17^{-/-}/Tg mice. The expression of other molecules (*Ccr6*, *Tnf*, *Il6*, *Il22*, *Il17f*, *Rorc*, *Tbx21*, and *Gata3*) was not suppressed in those mice (Fig. 5F). These results indicate that the production of IL-17 is not essential for the development of sialadenitis in ROR γ T Tg mice.

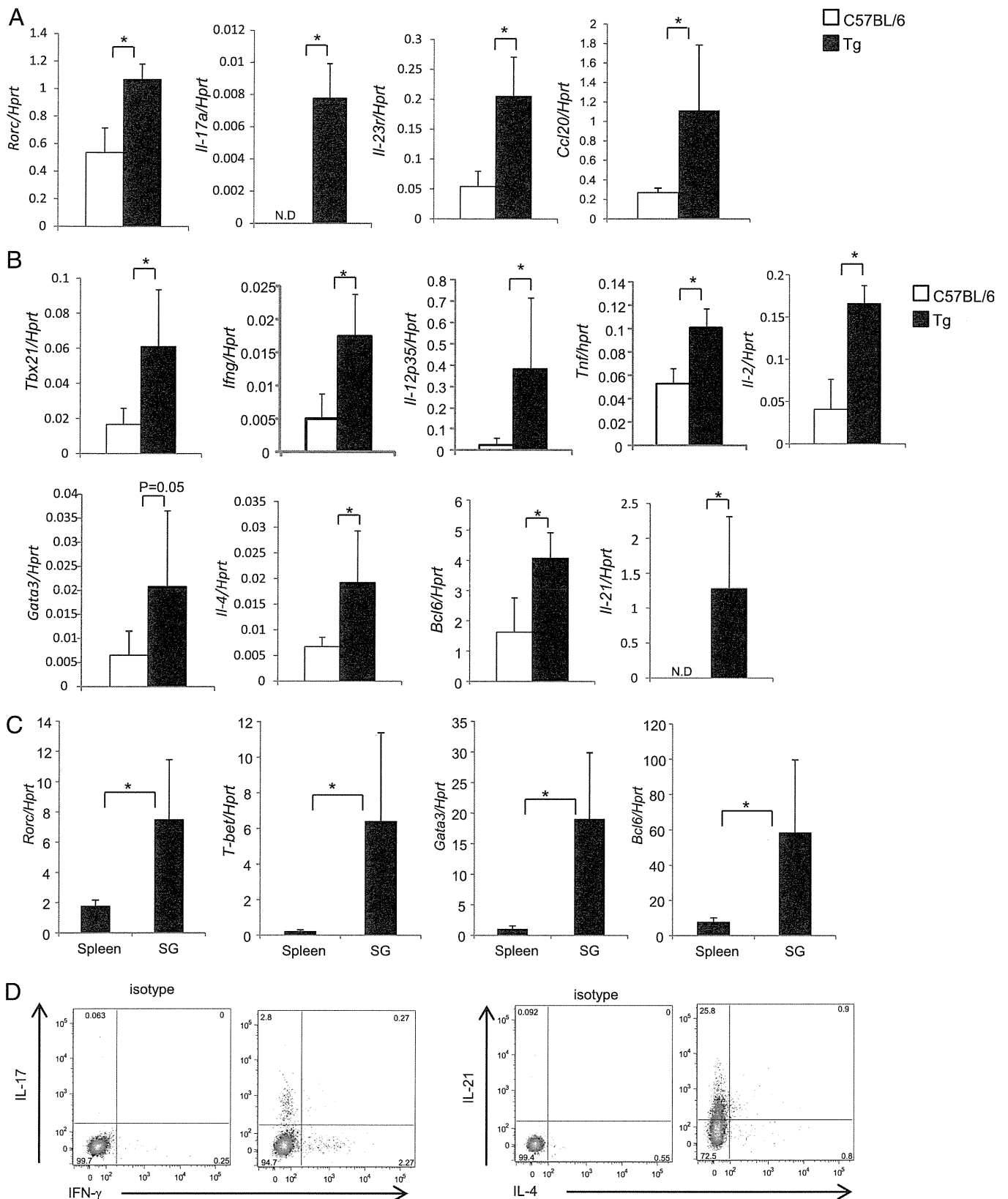


FIGURE 4. Detection of Th1, Th2, Th17, and Tfh cell-related factors in inflammatory lesions. **(A and B)** Quantitative PCR analysis of *Rorc*, *Il17a*, *Il21*, *Il23r*, *Ccl20*, *Ifng*, *Il2*, *Il12*, *Tnf*, *Il4*, *Tbx21*, *Gata3*, and *Bcl6* mRNA expression in salivary glands of C57BL/6 and ROR γ t Tg mice. The experiment was performed in duplicate. **(C)** Quantitative PCR analysis of mRNA expression in CD4⁺ T cells isolated from the spleen and salivary glands (SG) of ROR γ t Tg mice. Data were normalized to the expression of the reference gene, *Hprt*. **(D)** Flow cytometry analysis of infiltrating CD4⁺ T cells in salivary glands of ROR γ t Tg mice after stimulation with PMA and ionomycin for 4 h. Four mice per group were analyzed and representative data are shown. **p* < 0.05 (Mann-Whitney *U* test).

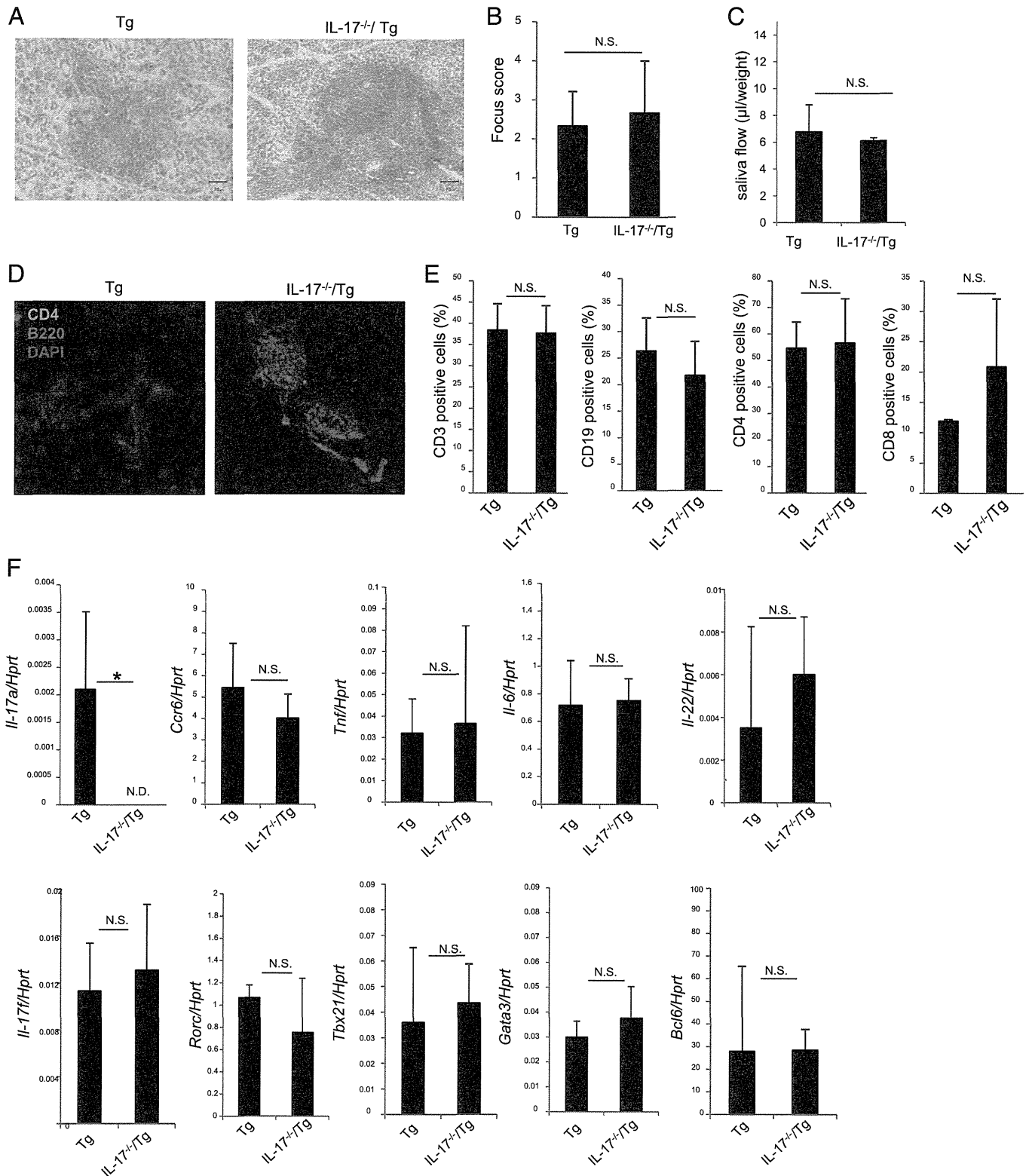


FIGURE 5. Spontaneous development of sialadenitis in IL-17^{-/-}/Tg mice. **(A)** H&E-stained sections of salivary glands from ROR γ t Tg and IL-17^{-/-}/ROR γ t Tg (IL-17^{-/-}/Tg) mice. Scale bars, 70 μ m. Data are representative from one of the four tissue samples with similar results. **(B)** Histological score of inflammatory lesions in the salivary glands of animals described in (A). **(C)** Saliva volume of ROR γ t Tg and IL-17^{-/-}/Tg mice. **(D)** Immunofluorescence analysis of salivary glands from ROR γ t Tg and IL-17^{-/-}/Tg. Data are representative of four tissue samples with similar results. Original magnification \times 100. **(E)** Flow cytometric analysis of lymphocyte gated cells (15–20%) stained with CD19, CD3, CD4, and CD8 on CD3⁺ subsets in infiltrated cells isolated from salivary glands of ROR γ t Tg and IL-17^{-/-}/Tg mice at 12 wk of age. Data are representative of two independent experiments with four mice per group. **(F)** Quantitative PCR analysis of mRNA expression level of *Il17a*, *Il17f*, *Ccr6*, *Tnf*, *Il6*, *Il22*, *Rorc*, *Tbx21*, and *Gata3* in salivary glands of ROR γ t Tg and IL-17^{-/-}/Tg mice. The experiment was performed in duplicate. Data were normalized to the expression of the reference gene, *Hprt*. * p < 0.05 (Mann–Whitney U test).

Reduced $CD4^+CD25^+Foxp3^+$ Treg cells in $ROR\gamma t$ Tg mice

Our data showed that IL-17 is not involved in the development of sialadenitis in $ROR\gamma t$ Tg mice (Fig. 5). We then examined whether overexpression of $ROR\gamma t$ resulted in inhibition of Foxp3 expression in $CD4^+CD25^+$ T cells, because $CD4^+CD25^+Foxp3^+$ Treg cells are known to inhibit onset of autoimmune disease. To answer this question, we first analyzed the expression of Foxp3 in $CD4^+CD25^+$ cells from $ROR\gamma t$ Tg mice. The expression of Foxp3 and the proportion of Treg cells were significantly lower in the thymus, spleen, and cLN of $ROR\gamma t$ Tg mice compared with C57BL/6 mice (Fig. 6A, 6B). Second, the expression of CD25, a Treg cell signature gene, was significantly lower in Treg cells of $ROR\gamma t$ Tg mice compared with that in the same cells of C57BL/6 mice. Alternatively, the staining intensities of CTLA4, GITR, and CD103 were higher in $ROR\gamma t$ Tg mice than in C57BL/6 mice (Fig. 6C, 6D), which was probably secondary to ongoing inflammation in $ROR\gamma t$ Tg mice. In Treg cells of $ROR\gamma t$ Tg mice, expression of $ROR\gamma t$ was also significantly higher than in those of C57BL/6 mice (Fig. 6C, 6D). Moreover, we examined the role of IL-2 signaling in downregulation of Foxp3 expression, because the above results showed downregulation of IL-2 receptor CD25 expression in Treg cells of $ROR\gamma t$ Tg mice. Phosphorylation of STAT5 was significantly lower and the expression level of *Socs1* mRNA was significantly increased in IL-2-stimulated Treg cells in $ROR\gamma t$ Tg mice compared with those in C57BL/6 mice (Fig. 6E, 6F). Alternatively, TGF- β -induced smad2/3 phosphorylation was not inhibited in Treg cells of $ROR\gamma t$ Tg mice (Supplemental Fig. 4A).

We also investigated the effect of $ROR\gamma t$ overexpression on the inhibitory function of Treg cells using Foxp3-GFP reporter mice (C57BL/6-Foxp3^{GFP}, $ROR\gamma t$ -Foxp3^{GFP} mice). The Treg cells are known to be unresponsive to TCR stimulation and do not produce effector cytokines in vitro (21). We found that $ROR\gamma t$ -overexpressing $CD4^+CD25^+GFP^+$ cells produced large amounts of IL-17 and IL-10 compared with C57BL/6-Foxp3^{GFP} mice (Fig. 6G). Moreover, we also tested the ability of Treg cells to suppress effector cell proliferation in vitro by comparing the function of $CD4^+CD25^+GFP^+$ cells isolated from $ROR\gamma t$ -Foxp3^{GFP}, C57BL/6-Foxp3^{GFP} mice in inhibiting the proliferation of $CD4^+CD25^-GFP^-$ cells from C57BL/6-Foxp3^{GFP} mice. The suppressive activity of $CD4^+CD25^+GFP^+$ cells isolated from $ROR\gamma t$ -Foxp3^{GFP} mice was equal to that of the same cells isolated from C57BL/6-Foxp3^{GFP} mice (Fig. 6H, 6I). In contrast, the suppressive activity of $CD4^+CD25^+GFP^-$ cells, which were significantly higher in $ROR\gamma t$ Tg mice, on effector T (Teff) cells was less than $CD4^+CD25^+GFP^+$ cells in $ROR\gamma t$ Tg mice (Supplemental Fig. 4B, 4C). These figures showed that $ROR\gamma t$ overexpression in Treg cells has no effect on regulatory function.

Development of sialadenitis via reduced Treg cells

To evaluate the effect of reduction of Treg cells on prevention of sialadenitis, we coinoculated $CD4^+CD25^-GFP^-$ cells (as Teff cells) and Treg cells isolated from C57BL/6-Foxp3^{GFP} or $ROR\gamma t$ -Foxp3^{GFP} mice into $Rag2^{-/-}$ mice. Such inoculation was associated with the lack of development of sialadenitis in Teff (C57BL/6)→ $Rag2^{-/-}$ mice, in contrast to the appearance of sialadenitis in Teff (Tg)→ $Rag2^{-/-}$ mice (Fig. 7A). Infiltrated cells were not detected in the salivary glands of Teff (Tg) + Treg (Tg, 5×10^5 cells)→ $Rag2^{-/-}$ mice, as well as in those of Teff (Tg) + Treg (C57BL/6, 5×10^5 cells)→ $Rag2^{-/-}$ mice, and the histological score of inflammatory lesions in the salivary glands of Teff (Tg) + Treg (Tg)→ $Rag2^{-/-}$ mice was similar to score of Teff (Tg) + Treg (C57BL/6)→ $Rag2^{-/-}$ mice (Fig. 7B). These data showed that abundant Treg cells from $ROR\gamma t$ Tg mice could inhibit the

development of sialadenitis. However, the reduced Treg cells could not inhibit sialadenitis in Teff (Tg) + 1:10 Treg (Tg, 5×10^4 cells)→ $Rag2^{-/-}$ mice (Fig. 7B). These findings indicated that Teff cells of $ROR\gamma t$ Tg mice was required for induction of sialadenitis, and the reduction of Treg cells in $ROR\gamma t$ Tg mice did not prevent the development of sialadenitis, suggesting its involvement in the pathogenesis of SS.

Discussion

Although IL-17 and IL-23 produced by Th17 cells are associated with the pathogenesis of SS, the pathological role of Th17 cells remains obscure (11–15). The results of the present study shed light on the function of $ROR\gamma t$ in the development of sialadenitis. Infiltrating mononuclear cells in the salivary glands of $ROR\gamma t$ Tg mice were mainly $CD4^+$ T cells at the early stage of the inflammation whereas B cells gradually increased with age. These changes resemble lymphocyte infiltration in the labial salivary glands of patients with SS. The results provide evidence for the role of $ROR\gamma t$ overexpression in $CD4^+$ T cells in the development of spontaneous sialadenitis-like SS through downregulation of $CD4^+CD25^+Foxp3^+$ Treg cells.

Experimental evidence suggests that IL-17 produced from $CD4^+$ T cells is directly involved in the onset of SS-like disease (11, 22). However, we demonstrated in the present study the development of sialadenitis in IL-17-deficient $ROR\gamma t$ Tg mice, indicating that IL-17 is not essential for the development of sialadenitis in $ROR\gamma t$ Tg mice. Our results showed expression of other inflammatory cytokines (*Tnf* and *Il6*) and Th1-, Th2-, and Tfh-related factors in the salivary glands of IL-17^{-/-}/Tg mice and $ROR\gamma t$ Tg mice. These results suggest that various cytokines are involved in the inflammation of sialadenitis in $ROR\gamma t$ Tg mice. Previous work provides evidence that although Th1 and Th2 effector memory cells could not be converted into Th17 cells, Th17 cells could be converted into Th1 and Th2 cells via the upregulation of *Tbet* and *Gata3* expression in presence of IL-12 or IL-4 (23, 24). We detected the higher expression of *Tbet*, *Gata3*, and *Bcl6* in $CD4^+$ T cells of salivary glands than those in splenic $CD4^+$ T cells of $ROR\gamma t$ Tg mice. Therefore, $ROR\gamma t$ -overexpressed $CD4^+$ T cells might convert into various T cells subsets (Th1, Th2, and Tfh) in salivary glands despite having the Th17-like phenotype in lymphoid tissue.

In $CD4^+$ T cells of $ROR\gamma t$ Tg mice, high expression of *Ccr6*, which is induced by $ROR\gamma t$ (17, 25), and *Ccl20* in the salivary glands of $ROR\gamma t$ Tg mice was noted, which might be involved in migration of $ROR\gamma t$ highly expressing $CD4^+$ T cells into salivary glands. The high expression of *Ccl20* in the salivary glands of patients with SS may be triggered by previous bacterial or viral infections in the local regions (26–29).

$ROR\gamma t$ is also crucial role for the development and function of lymphoid tissue inducer (LTi) cells, which share proinflammatory cytokines with Th17 cells, as well as their requirement for $ROR\gamma t$ (30). $CD4^+$ LTi cells from spleen of postnatal mice express both IL-17 and IL-22 (31). Because IL-17 was not involved in the generation of sialadenitis, and $CD4^+$ LTi cells were not detected in the inflammatory lesions of $ROR\gamma t$ Tg mice (data not shown), $CD4^+$ LTi cells may not engage in the pathogenesis of sialadenitis in our model. However, the expression of *Il22* was not suppressed in salivary glands of IL-17^{-/-}/Tg mice. Further analysis of participation of IL-22 producer cells (including Th17, LTi cells) in the sialadenitis is needed.

Our findings showed the presence of activated $CD4^+$ T cells and $CD4^+CD25^+Foxp3^+$ cells that expressed CTLA4, GITR, and CD103. These findings are thought to be related to dysfunctional

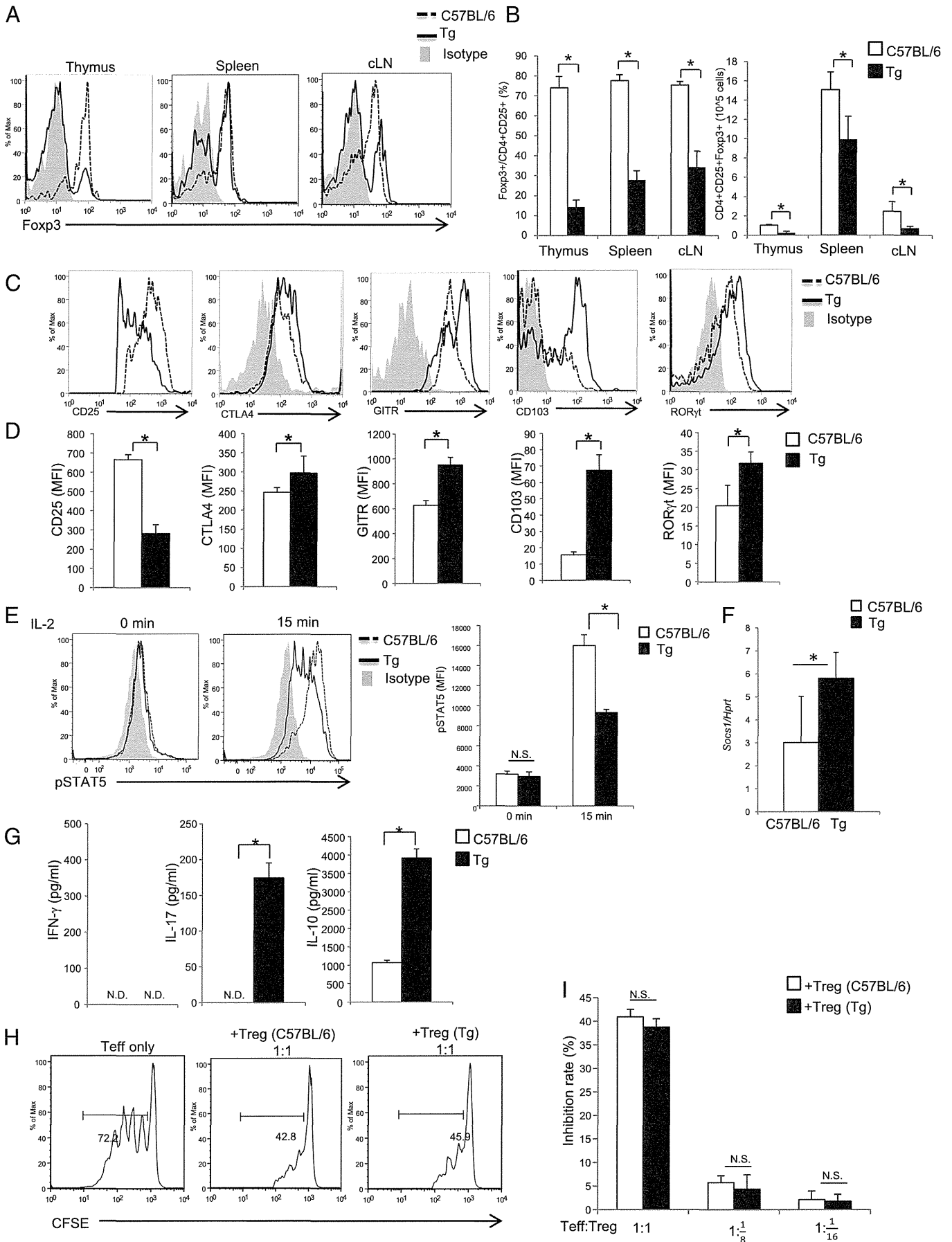


FIGURE 6. Downregulation of Foxp3 expression in CD4⁺CD25⁺ Treg cells. **(A)** Flow cytometric analysis of Foxp3 expression in CD4⁺CD25⁺ cells in thymus, spleen, and cLN of C57BL/6 and ROR γ t Tg mice. Data are representative of two independent experiments with four mice per group. **(B)** Proportion of Foxp3⁺ cells gated on CD4⁺CD25⁺ and CD4⁺CD25⁺Foxp3⁺ cells in thymus, spleen, and cLN of C57BL/6 and ROR γ t (Figure legend continues)

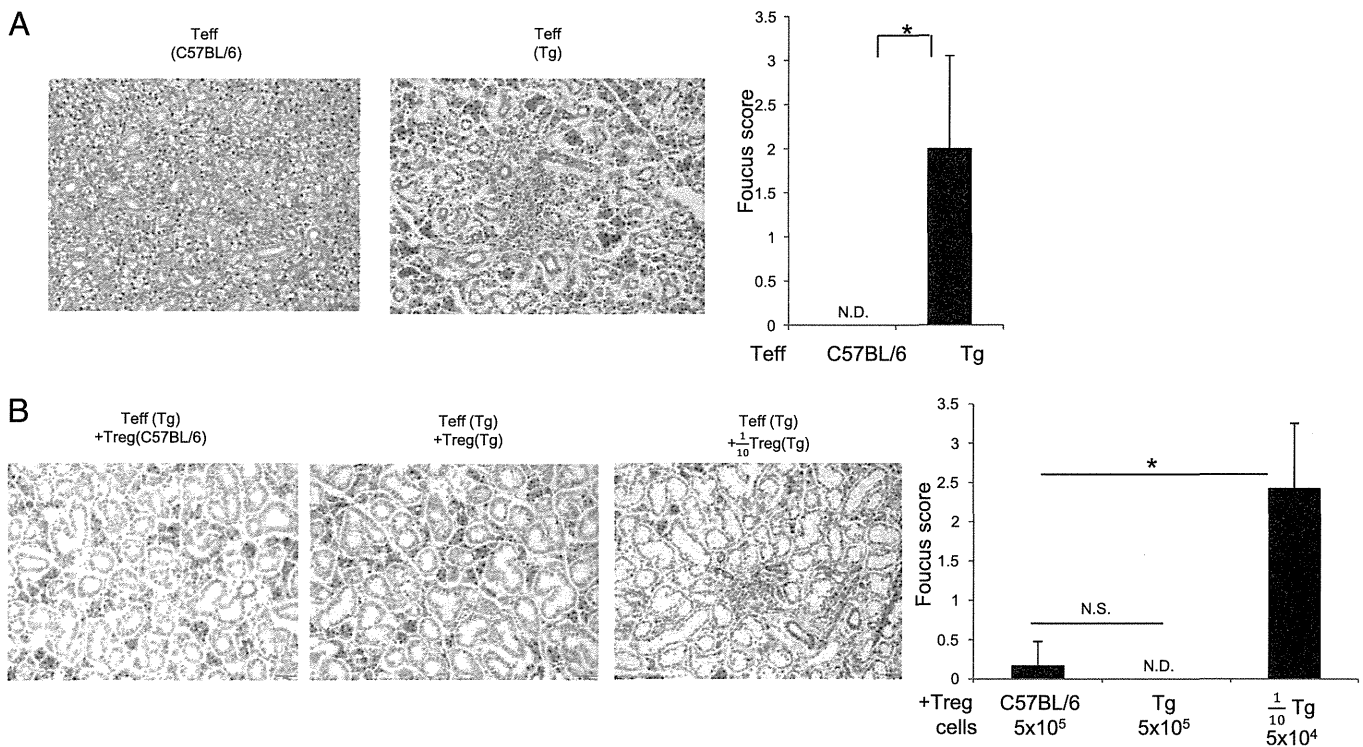


FIGURE 7. Function of ROR γ t-overexpressing Treg cells in inhibition of sialadenitis. **(A)** CD4⁺CD25⁻GFP⁻ cells (Teff) (2×10^6 cells) isolated from C57BL/6-Foxp3^{GFP} or ROR γ t Tg-Foxp3^{GFP} mice were inoculated into Rag2^{-/-} mice. * $p < 0.05$ (Mann-Whitney U test). **(B)** CD4⁺CD25⁻GFP⁻ cells (Teff) (2×10^6 cells) isolated from ROR γ t Tg-Foxp3^{GFP} mice were inoculated into Rag2^{-/-} mice with CD4⁺CD25⁺GFP⁺ cells (Treg) (5×10^5 cells) of C57BL/6-Foxp3^{GFP} mice or CD4⁺CD25⁺GFP⁺ cells (Treg) (5×10^5 or 5×10^4 [1:10] cells) of ROR γ t Tg-Foxp3^{GFP} mice. The bar graph shows the histological score of inflammatory lesions in the salivary glands. At 22 wk postinoculation, the pathology of salivary glands was analyzed. Salivary glands were sectioned at 4 μ m, and each section was stained with H&E. Scale bars, 50 μ m. Data are representative of four tissue samples with similar results. Histological score of inflammatory lesions in the salivary glands of the experiment described. * $p < 0.05$ (Tukey t test).

Treg cells (32). Based on this finding, we hypothesized that ROR γ t overexpression inhibits the expression of Foxp3 in CD4⁺CD25⁺ cells. Foxp3 can suppress Th17 cell differentiation through antagonism of ROR γ t activity (33), and its interaction has been reported (34). Our results showed that the expression of Foxp3 was significantly decreased in CD4⁺CD25⁺ cells, indicating that downregulation of Foxp3 expression in these cells is the culprit mechanism in sialadenitis. Deficiency of Treg cells is probably one of the underlying mechanisms of SS, because massive leukocyte infiltration was observed in the salivary glands of IL-2R α ^{-/-}, IL-2^{-/-}, and Rag1^{-/-} mice inoculated with lymph node cells from scurfy mice and other mice with impaired Treg function (32, 35, 36). In adoptive transfer models, such as CD4⁺CD45RB^{hi} naive T cells without Treg cells in Rag-deficient mice, sialadenitis was also observed (37). Previous studies demonstrated the presence of significantly lower CD4⁺CD25⁺ Treg cell count in peripheral blood of patients with SS compared with healthy controls. However, the

inhibitory function of these cells in SS was not different from that of the control (39). Recent studies have reported a close correlation between the proportion of Foxp3⁺ Treg cells in the salivary glands of patients with SS and the severity of inflammation as well as certain risk factors for lymphoma development (39, 40). In the present study, suppressive activity of Treg cells in ROR γ t Tg mice was equal to that of the same cells in C57BL/6 mice, and sialadenitis was not observed in Rag2^{-/-} mice inoculated with effector cells plus Treg cells from ROR γ t Tg mice. However, we found sialadenitis in Rag2^{-/-} mice inoculated with effector cells plus decreased numbers of Treg cells from ROR γ t Tg mice. These data provide strong support to the notion that the reduction of Treg cells in ROR γ t Tg mice might play a role in the development of sialadenitis.

How can overexpression of ROR γ t result in suppression of Foxp3 expression? Judging from the low level of CD25 expression and the inhibition of IL-2-STAT5 signaling in Treg cells from

Tg mice. Four mice per group were analyzed and representative data of two independent experiments with consistent results are shown. **(C)** Flow cytometric analysis of CD25, CTLA4, GITR, CD103, and ROR γ t in splenocytes from C57BL/6 and ROR γ t Tg mice. These cells were gated on CD4⁺CD25⁺Foxp3⁺ cells. Data are representative of two independent experiments with four mice per group. **(D)** Mean fluorescence intensity (MFI) of CD4⁺CD25⁺Foxp3⁺ cells. Data are representative of two independent experiments with four mice per group. **(E)** Detection of phosphorylation of STAT5 after stimulation with 1 μ g/ml IL-2 in CD4⁺CD25⁺Foxp3⁺ cells from C57BL/6-Foxp3^{GFP} or ROR γ t Tg-Foxp3^{GFP} mice by flow cytometry. The bar graph shows the MFI in CD4⁺CD25⁺Foxp3⁺ cells. Data are representative of two independent experiments with four mice per group. **(F)** Quantitative PCR analysis of mRNA expression level of *Socs1* in IL-2-stimulated Treg cells of C57BL/6 and ROR γ t Tg mice. The experiment was performed in duplicate. Data were normalized to the expression of the reference gene, *Hprt*. **(G)** Cytokine secretion (IFN- γ , IL-17, IL-10) in CD4⁺CD25⁺Foxp3⁺ cells from C57BL/6-Foxp3^{GFP} or ROR γ t Tg-Foxp3^{GFP} mice stimulated for 72 h with anti-CD3/28 beads. Data are representative of three independent experiments with four mice per group. **(H)** CFSE-labeled CD4⁺CD25⁺Foxp3⁻ cells from C57BL/6-Foxp3^{GFP} mice cultured with or without CD4⁺CD25⁺Foxp3⁺ cells from C57BL/6-Foxp3^{GFP} or ROR γ t Tg-Foxp3^{GFP} mice and stimulated with anti-CD3/28 beads for 72 h. Data are CFSE gated and representative of three independent experiments with four mice per group. **(I)** The bar graph shows the rate of inhibition. * $p < 0.05$ (Mann-Whitney U test). N.D., detected.

ROR γ Tg mice, we suggest that the impaired IL-2 signaling in Treg cells could be involved in the downregulation of Foxp3 expression. Expression of CD25 is directly linked to IL-2 signaling, and its signaling is important for maintaining homeostasis of Treg cells in vivo (41). Previous studies indicated that IL-2R–dependent STAT5 activation directly drives the development of Treg cells, which includes maturation of immature Treg cells into mature CD4⁺CD25⁺Foxp3⁺ Treg cells in the thymus. Furthermore, STAT5 binds to the conserved sites within the Foxp3 promoter, and then STAT5 signaling regulates Foxp3 expression (42, 43). Our data showed that *Socs1* expression in IL-2–stimulated Treg cells of ROR γ Tg mice was significantly higher than in that of C57BL/6 mice. SOCS1 was the protein with a known inhibitory role in the IL-2 signaling pathway leading to reduced STAT5 phosphorylation (44). STAT5 was reported to be phosphorylated in SOCS1-deficient Treg cells at higher levels than in control mice, which is associated with the increase in number of Treg cells (45). In Treg cells from ROR γ Tg mice, SOCS1 might inhibit the STAT5 phosphorylation after stimulation with IL-2. Mice deficient in IL-2 and CD25 are known to develop systemic autoimmune disease due to impaired development and function of Treg cells (46, 47). It has been reported that ROR γ T negatively regulates IL-2 production in T cells (48, 49).

The conserved noncoding sequence (CNS) has been characterized as regulatory regions controlling induction and maintenance of Foxp3 expression. CNS1 was regulated by Smad3 and NFAT, and naive CD4⁺ T cells from CNS1-lacking mice displayed reduced capacity to upregulate Foxp3 expression in the presence of TGF- β (50). CNS2 was also crucial role for the maintenance of stable level of Foxp3 expression, which was controlled by STAT5-regulated TGF- β signaling through Smad3 (51). The factor of TGF- β in addition to IL-2 is necessary for the development of Treg cells. In ROR γ Tg mice, TGF- β –induced Smad signaling was not inhibited in Treg cells, and hence disruption of Smad3-mediated TGF- β signaling did not involve the downregulation of Foxp3 expression.

Our results demonstrated that both ROR γ T-overexpressed CD4⁺ T cells and reduced Treg cells via suppression of IL-2–induced phosphorylation of STAT5 might contribute to the development of SS-like sialadenitis. These results suggest that regulation of ROR γ T expression is a potentially useful therapeutic strategy against SS.

Disclosures

The authors have no financial conflicts of interest.

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TGF- β 3-expressing CD4⁺CD25⁻LAG3⁺ regulatory T cells control humoral immune responses

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Autoantibodies induce various autoimmune diseases, including systemic lupus erythematosus (SLE). We previously described that CD4⁺CD25⁻LAG3⁺ regulatory T cells (LAG3⁺ Treg) are regulated by Egr2, a zinc-finger transcription factor required for the induction of T-cell anergy. We herein demonstrate that LAG3⁺ Treg produce high amounts of TGF- β 3 in an Egr2- and Fas-dependent manner. LAG3⁺ Treg require TGF- β 3 to suppress B-cell responses in a murine model of lupus. Moreover, TGF- β 3- and LAG3⁺ Treg-mediated suppression requires PD-1 expression on B cells. We also show that TGF- β 3-expressing human LAG3⁺ Treg suppress antibody production and that SLE patients exhibit decreased frequencies of LAG3⁺ Treg. These results clarify the mechanism of B-cell regulation and suggest therapeutic strategies.

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Autoantibodies induce various autoimmune diseases, including systemic lupus erythematosus (SLE)¹, which is characterized by severe inflammation in multiple organ systems. The high-affinity autoantibodies primarily originating from the self-reactive B cells underwent somatic hypermutation in the germinal centre (GC)². Follicular helper T (T_{FH}) cells expressing CXCR5 have emerged as a lineage of helper T cells (Th cells) that are functionally specialized to provide help to B cells, allowing the formation of GC and the subsequent long-lived plasma cell differentiation. Therefore, regulation of the quality and quantity of T_{FH} cells and memory B-cell populations in GC (GCB) is important to prevent immunopathology. CD4⁺CD25⁺Treg (CD25⁺Treg) that express Foxp3 play the key roles in the maintenance of self-tolerance and suppress the activation of conventional T cells and dendritic cells³. Moreover, accumulating evidence indicates the essential role of CD25⁺Treg, including CD4⁺CD25⁺CXCR5⁺ follicular Treg² and CD4⁺CD25⁺CD69⁻Treg⁴, in the regulation of humoral immunity. These observations highlight the protective role of CD25⁺Treg in systemic autoimmunity; however, the disease induced by the absence of functional CD25⁺Treg is quite different from SLE^{1,5}. Moreover, a role for CD25⁺Treg in SLE has not been clearly established⁶. Recent advances in understanding of CD8⁺Treg have underscored the importance of Qa-1-restricted CD8⁺Treg for the maintenance of B-cell tolerance. Mice with functional impairment in CD8⁺Treg exhibit a lupus-like disease with a significant increase in T_{FH}⁷. The development of systemic autoimmunity in B6.Yaa mutant mice is associated with a pronounced defect in CD8⁺Treg activity⁸. Nevertheless, the actual contribution of CD8⁺Treg to the regulation of human autoimmunity remains unclear.

Early growth response gene 2 (Egr2), a zinc-finger transcription factor, plays a critical role in hindbrain development and myelination of the peripheral nervous system⁹. In T cells, Egr2 is important for the maintenance of T-cell anergy by negatively regulating T-cell activation¹⁰. The involvement of Egr2 in the control of systemic autoimmunity was first suggested by the observation that lymphocyte-specific Egr2-deficient mice develop a lupus-like disease with no impact on the development of Foxp3-expressing CD25⁺Treg¹¹. Moreover, mice deficient for both Egr2 and Egr3 in B and T cells present lethal and early-onset systemic autoimmunity, suggesting a synergistic role for Egr2 and Egr3 in controlling B-cell tolerance¹². We and our collaborators have shown that polymorphisms in *EGR2* influence SLE susceptibility in humans¹³. We have previously identified Egr2-controlled CD4⁺CD25⁻LAG3⁺Treg (LAG3⁺Treg)¹⁴. LAG3 is a CD4-related molecule that binds to MHC class II, and the binding induces immunoreceptor tyrosine-based activation motif (ITAM)-mediated inhibitory signalling¹⁵. Approximately 2% of the CD4⁺CD25⁻T-cell population in the spleen express LAG3. These LAG3⁺Treg produce high levels of interleukin (IL)-10 and are suppressive in a murine model of colitis in an IL-10-dependent manner. Unlike CD25⁺Treg, high-affinity interactions with selecting peptide/MHC ligands expressed in the thymus do not induce the development of LAG3⁺Treg. Recently, Gagliani *et al.*¹⁶ reported that concomitant expression of LAG3 and CD49b is specific for IL-10-producing type 1T regulatory (Tr1) cells, confirming that LAG3 is one of the phenotypic markers of IL-10-producing Foxp3-independent CD4⁺Treg.

The association between Egr2 and autoantibody-mediated systemic autoimmunity suggested a linkage between Egr2-expressing LAG3⁺Treg and the control of B-cell responses. We herein demonstrate that LAG3⁺Treg produce high amounts of transforming growth factor-β3 (TGF-β3) and suppress B-cell development and antibody production. In

MRL/lpr lupus-prone mice, adoptive transfer of LAG3⁺Treg from MRL/+ mice suppresses the progression of lupus in a TGF-β3-dependent manner. Expression of both Fas and Egr2 by LAG3⁺Treg is necessary for TGF-β3 production and for the suppression of humoral immunity. These results clarify the mechanisms underlying LAG3⁺Treg-mediated B-cell regulation.

Results

Egr2 mediates control of humoral immunity by LAG3⁺Treg.

To clarify the role of Egr2 in T cells, we generated T-cell-specific Egr2 conditional knockout (CKO) mice (*Egr2^{fl/fl}* CD4-*Cre*⁺). Egr2 CKO mice showed significant increases in the proportion of CD4⁺CD25⁻CXCR5⁺PD-1⁺T_{FH} and B220⁺GL-7⁺Fas⁺GCB (Fig. 1a), and they demonstrated enhanced 4-hydroxy-3-nitrophenylacetyl (NP)-specific antibody production following a single immunization with NP-ovalbumin (NP-OVA; Fig. 1b and Supplementary Fig. 1a). Transfer of wild-type (WT) LAG3⁺Treg significantly suppressed the spontaneous differentiation of T_{FH} and GCB (Fig. 1a) and inhibited excessive antibody production (Fig. 1b), indicating that LAG3⁺Treg are able to suppress B-cell responses *in vivo*. To examine whether physiological number of WT LAG3⁺Treg could improve excessive T_{FH} and GCB development in Egr2 CKO mice, we reconstituted Thy1.2⁺Egr2 CKO-recipient mice with an equal number of bone marrow (BM) cells derived from Thy1.2⁺Egr2 CKO mice and Thy1.1⁺WT mice (Supplementary Fig. 2). Six weeks after the reconstitution, although less than 50% of splenocytes were derived from Egr2-sufficient WT mice, the excessive development of T_{FH} and GCB cells was significantly reduced (Supplementary Fig. 2b,c). This result suggested that the immunological abnormality of Egr2 CKO mice is dependent, at least in part, on the defects in Egr2-expressing LAG3⁺Treg. In an *in vitro* T-cell/B-cell co-culture system, anti-CD3-stimulated WT LAG3⁺Treg more efficiently reduced the percentage of viable anti-IgM-stimulated B cells as well as total IgG production from anti-CD40/IL-4-stimulated B cells when compared with CD25⁺Treg (Fig. 1c,d). To evaluate B-cell responses *in vivo*, recombination-activating gene 1-deficient (Rag1KO) mice were transferred with WT B cells and CD4⁺CD25⁻LAG3⁻ helper T cells (Th cells) from OVA-specific OT-II T cell receptor (TCR) transgenic mice, and then immunized with NP-OVA twice. Strikingly, co-transfer of WT LAG3⁺Treg effectively suppressed NP-specific antibody responses and the development of T_{FH} and GCB (Fig. 1e,f and Supplementary Fig. 1b). The enhanced GCB development and antibody production in Egr2 CKO mice suggested a pivotal role for Egr2 in B-cell regulation. Although Egr2 confers the phenotype of LAG3⁺Treg¹⁴, T cells with the phenotype of LAG3⁺Treg were also observed in the spleen of Egr2 CKO mice (Supplementary Fig. 1c). This result suggested that the expression of LAG3, the marker for LAG3⁺Treg, is not fully dependent on Egr2. Egr2-deficient LAG3⁺Treg (Supplementary Fig. 1c) failed to suppress *in vivo* B-cell antibody production and the development of T_{FH} and GCB (Fig. 1e,f). Thus, the expression of Egr2 on LAG3⁺Treg is necessary for the suppression of B-cell responses. In transgenic mice that express green fluorescent protein (GFP) under the control of the Egr2 promoter (Egr2-GFP mice; Supplementary Fig. 3a), the expression of GFP in CD4⁺T cells correlated with Egr2 protein expression (Supplementary Fig. 3b). The importance of Egr2 was confirmed by the observation that CD4⁺CD25⁻Egr2-GFP⁺ cells from Egr2-GFP mice also exhibited B-cell-suppressive activity *in vivo*, similar to that of LAG3⁺Treg (Supplementary Fig. 3c). We next determined whether the suppression of antibody production via LAG3⁺Treg is induced not only under lymphopenic conditions, such as Rag1KO mice, but also under more physiological non-

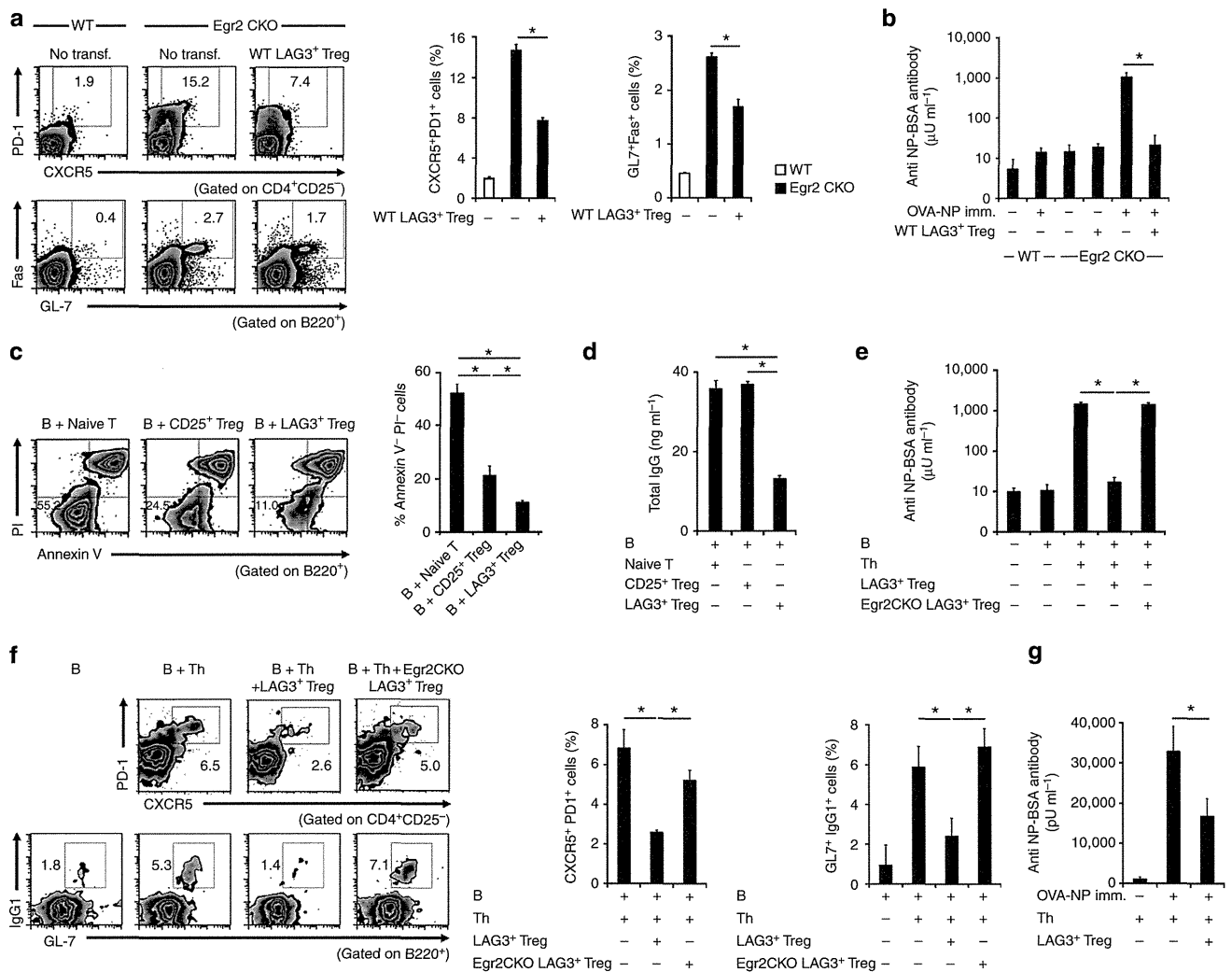


Figure 1 | LAG3⁺ Treg exhibit Egr2-dependent control of antibody production. (a) Flow cytometry plots and quantification of CD4⁺CD25⁻CXCR5⁺PD-1⁺ T_{FH} and B220⁺GL-7⁺Fas⁺ GCB from WT or Egr2 CKO mice at 7 days with or without the adoptive transfer of WT LAG3⁺ Treg (n = 5 per group). *P < 0.05 (unpaired two-tailed Student's *t*-test). (b) NP-specific antibody responses of WT and Egr2 CKO mice immunized once with 100 μg NP-OVA/alum with or without adoptive transfer of WT LAG3⁺ Treg. The serum levels of anti-NP-BSA antibodies were analysed with ELISA 7 days after immunization. See also Supplementary Fig. 1a (n = 6 per group). *P < 0.05 (Bonferroni post-test). (c,d) *In vitro* B-cell suppression by LAG3⁺ Treg. Each T-cell subset stimulated with anti-CD3 mAb was co-cultured with stimulated B cells. (c) Live B220⁺ B cells were quantified with AnnexinV/PI staining 72 h after anti-IgM stimulation (n = 3 per group). *P < 0.05 (Bonferroni post-test). (d) Total IgM was determined in anti-CD40/IL-4-stimulated B-cell culture supernatants on day 7 by ELISA (n = 3 per group). *P < 0.05 (Bonferroni post-test). (e) *In vivo* NP-specific antibody responses. C57BL/6 (B6) B cells and OT-II CD4⁺CD25⁻LAG3⁻ Th cells were injected into Rag1KO mice in combination with or without LAG3⁺ Treg from B6 mice 1 day before the immunization with NP-OVA/alum, and given a booster immunization 14 days after the primary immunization. Anti-NP-BSA antibodies in sera were analysed with ELISA 7 days after the booster immunization. See also Supplementary Fig. 1b (n = 6 per group). *P < 0.05 (Bonferroni post-test). (f) Flow cytometry plots and quantification of splenic T_{FH} and B220⁺GL7⁺IgG1⁺ GCB from the same mice as in e. *P < 0.05 (Bonferroni post-test). (g) B-cell suppression by LAG3⁺ Treg in non-lymphopenic TEa mice. LAG3⁺ Treg from B6 mice and OT-II Th cells were injected into TEa mice and subsequently immunized with NP-OVA/alum once. Anti-NP-BSA antibody levels were determined with ELISA. See also Supplementary Fig. 1d (n = 6 per group). *P < 0.05 (post-test). Data are representative of three independent experiments. The means ± s.d. are indicated.

lymphopenic conditions. Ex peptide-specific TCR transgenic TEa mice were adoptively transferred with WT B cells and OT-II Th cells and subsequently immunized with NP-OVA once. Co-transferring WT LAG3⁺ Treg effectively suppressed NP-specific antibody production in non-lymphopenic TEa mice (Fig. 1g and Supplementary Fig. 1d). Next, the localization of LAG3⁺ Treg and CD25⁺ Treg was evaluated using Egr2-GFP mice and Foxp3-GFP mice, respectively. As shown in Supplementary Fig. 4a,b, Egr2-GFP⁺CD4⁺ cells were enriched in the T-B-cell border. In contrast, most of Foxp3-GFP⁺CD4⁺ cells were

located in the T-cell area. The difference of localization between LAG3⁺ Treg and CD25⁺ Treg may be associated with the functional variation between these regulatory subsets.

LAG3⁺ Treg suppress a lupus-like disease in MRL/lpr mice. We investigated whether LAG3⁺ Treg were able to inhibit disease progression in lupus-prone MRL-*Fas*^{lpr/lpr} (MRL/lpr) mice with a *Fas* mutation¹⁷. MRL/lpr mice were adoptively transferred with one of the various T-cell subsets from Fas-sufficient MRL-

Fas^{+/-} (MRL/+) mice. LAG3⁺ Treg, but not CD25⁺ Treg, significantly delayed proteinuria progression (Fig. 2a). Furthermore, the three-time transfers of LAG3⁺ Treg almost completely suppressed proteinuria progression. Increases in anti-ds DNA antibody titres and glomerular pathology scores were also inhibited by the single transfer of LAG3⁺ Treg (Fig. 2b–d and Supplementary Fig. 5a). In contrast, consistent with previous reports¹⁸, the three-time transfers of CD25⁺ Treg from MRL/+ mice to MRL/*lpr* mice did not alter the disease progression (Supplementary Fig. 5b–d). Furthermore, adoptive transfer of LAG3⁺ Treg from MRL/*lpr* mice had no therapeutic benefit in MRL/*lpr* mice (Supplementary Fig. 5b–d). Three-time injections of MRL/+ LAG3⁺ Treg also ameliorated lupus pathologies in MRL/*lpr* mice after the onset of overt proteinuria (Supplementary Fig. 5e,f).

The therapeutic effect of Fas-sufficient MRL/+ LAG3⁺ Treg in Fas-mutated MRL/*lpr* mice suggested that Fas contributes to the suppressive ability of LAG3⁺ Treg. Adding anti-FasL blocking antibody abrogated LAG3⁺ Treg-mediated antibody suppression both *in vitro* (Fig. 2e and Supplementary Fig. 1e) and *in vivo* (Fig. 2f–i). B6/*lpr* LAG3⁺ Treg, but not B6/*gld* LAG3⁺ Treg, failed to suppress antibody production (Fig. 2f–i). Therefore, Fas, but not FasL, on LAG3⁺ Treg is required to suppress B cells. Fas expression on CD4⁺ T cells was independent of *Egr2* because activated CD4⁺ T cells from WT and *Egr2* CKO mice expressed similar levels of Fas on LAG3⁺ cells (Supplementary Fig. 6).

TGF- β 3 produced by LAG3⁺ Treg controls humoral immunity. We next examined whether B-cell suppression by LAG3⁺

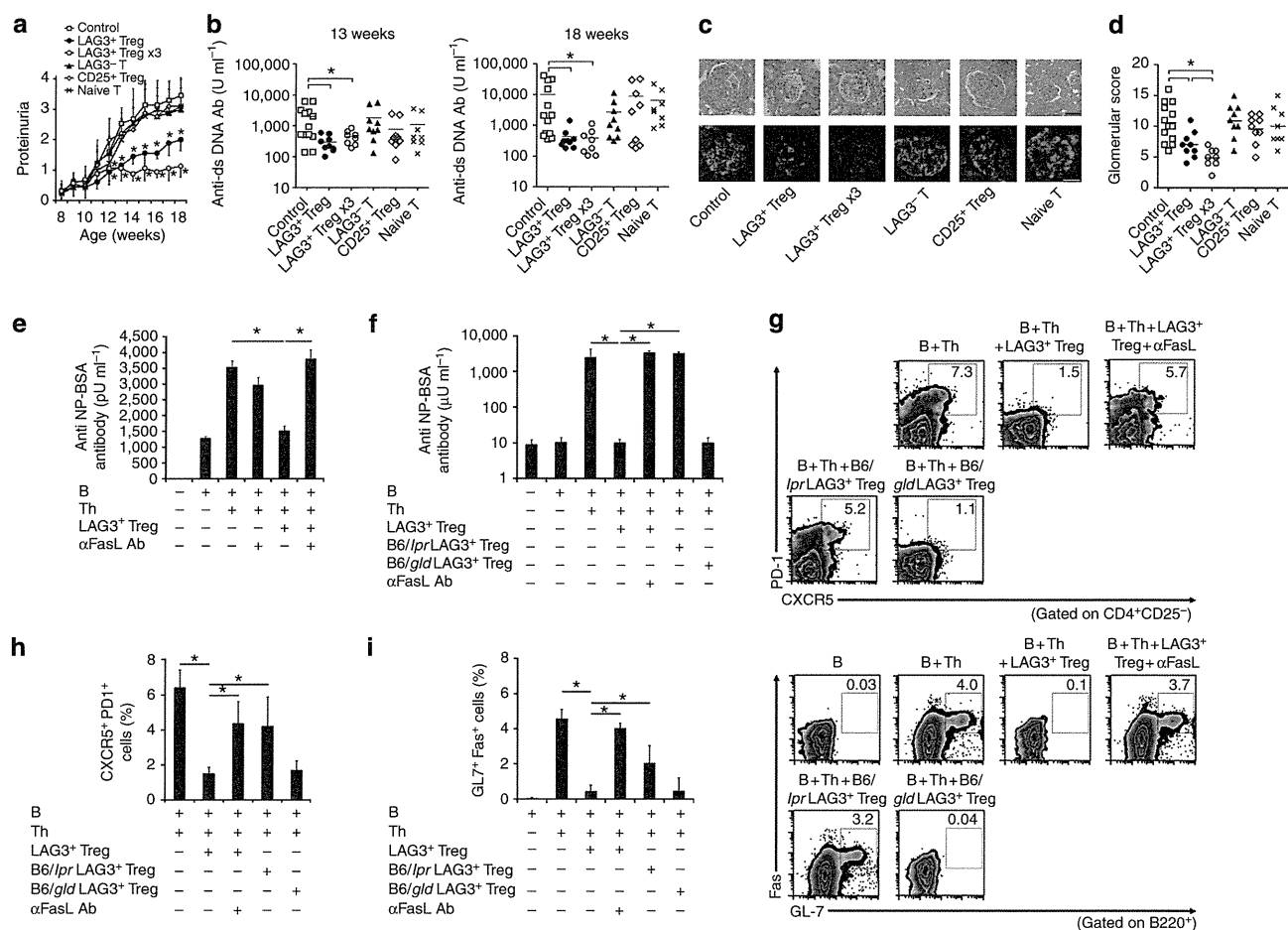


Figure 2 | LAG3⁺ Treg regulate B-cell functions through Fas. (a–d) Treatment of MRL/*lpr* mice with adoptive transfer of T-cell subsets. Ten-week-old MRL/*lpr* mice were injected i.v. with LAG3⁺ Treg ($n=9$), CD4⁺CD25⁻LAG3⁻ T cells (LAG3⁻ T; $n=9$), CD4⁺CD25⁺ Treg (CD25⁺ Treg; $n=9$) or CD4⁺CD25⁻CD45RB^{high} T cells (naive T; $n=8$) from MRL/+ mice (1×10^5 cells each). The mice of LAG3⁺ Treg x3 group ($n=8$) were injected with LAG3⁺ Treg (1×10^5 cells) at 10 weeks of age followed by a twice weekly injection of the same amount of LAG3⁺ Treg. The control group received PBS ($n=13$). (a) Proteinuria progression. * $P < 0.05$ versus control group (Mann-Whitney *U*-test). (b) Quantification of serum anti-ds DNA antibodies. * $P < 0.05$ (Bonferroni post-test). (c) Haematoxylin and eosine (H&E) staining (upper panels) and IgG immunofluorescent staining (lower panels) of kidney sections. Scale bars, 50 μ m. (d) Glomerular scores. * $P < 0.05$ (Mann-Whitney *U*-test). (e) LAG3⁺ Treg-mediated suppression of *in vitro* NP-specific antibody responses. B cells and Th cells purified from NP-OVA/alum-pre-immunized B6 mice and OT-II mice, respectively, were incubated with or without LAG3⁺ Treg from non-immunized OT-II mice in the presence or absence of anti-FasL blocking antibody, and supernatants were analysed for anti-NP-BSA antibodies using ELISA. See also Supplementary Fig. 1e ($n=6$ per group). * $P < 0.05$ (Bonferroni post-test). (f) NP-specific antibody responses of Rag1KO mice injected with B6 B cells and OT-II Th cells with or without LAG3⁺ Treg from WT, B6/*lpr* or B6/*gld* mice, as outlined in Fig. 1e ($n=6$ per group). Anti-FasL blocking antibody (200 μ g per mouse) was injected i.v. weekly. * $P < 0.05$ (Bonferroni post-test). (g–i) Flow cytometry plots (g) and quantification of splenic CD4⁺CD25⁻CXCR5⁺PD-1⁺ T_{FH} (h) and B220⁺GL7⁺Fas⁺ GCB (i) from the same mice as in f. Statistical significances in h,i were analysed by Bonferroni post-test (* $P < 0.05$). The experiments in e,f were repeated three times. The means \pm s.d. are indicated.

Treg is mediated by IL-10 or TGF- β family members. As described above, LAG3 is considered to be one of the specific cell-surface markers for Tr1 cells¹⁶. We previously reported that LAG3⁺ Treg produce large amounts of IL-10 (ref. 14), and Egr2 mediates IL-27-induced IL-10 production in CD4⁺ T cells through B-lymphocyte-induced maturation protein-1 (Blimp-1; coded by the *Prdm1* gene)¹⁹. As expected, IL-10 expression levels were significantly reduced in LAG3⁺ Treg from T-cell-specific Prdm1 CKO mice (*Prdm1^{fl/fl}* CD4-*Cre*⁺) compared with WT mice (Supplementary Fig. 7a). Although LAG3⁺ Treg derived from Prdm1 CKO and IL-10-deficient (IL-10KO) mice showed a slight reduction in the suppressive activity for *in vivo* NP-specific antibody responses compared with WT LAG3⁺ Treg, there were no statistical differences in suppressive activity among these three LAG3⁺ Treg (Supplementary Fig. 7b). These results suggested that IL-10 is not critical for B-cell suppression by LAG3⁺ Treg.

Microarray analysis¹⁴ and quantitative real-time PCR of LAG3⁺ Treg revealed a significant increase in TGF- β 3 expression, but not TGF- β 1 or 2 (Fig. 3a,b). TCR stimulation induced the production of a large amount of TGF- β 3, but not TGF- β 1 or 2, in the culture supernatants of LAG3⁺ Treg (Fig. 3c–e). The same trend was observed for the differences in mRNA levels of *Tgfb* families in LAG3⁺ Treg (Supplementary Fig. 8). In contrast, CD25⁺ Treg only produced small amounts of TGF- β 1 under these conditions. TGF- β 3 markedly suppressed anti-IgM-stimulated B-cell proliferation and CD40 expression (Fig. 3f), strongly induced B-cell death (Fig. 3g), and suppressed total IgG production (Fig. 3h). TGF- β 3 produced similar effects as TGF- β 1 and 2 (Supplementary Fig. 9a,b), in accordance with previous findings that TGF- β 1 strongly suppresses B-cell functions²⁰. Regarding signal transduction, the addition of TGF- β 3 significantly reduced the phosphorylation of signal transducer and activator of

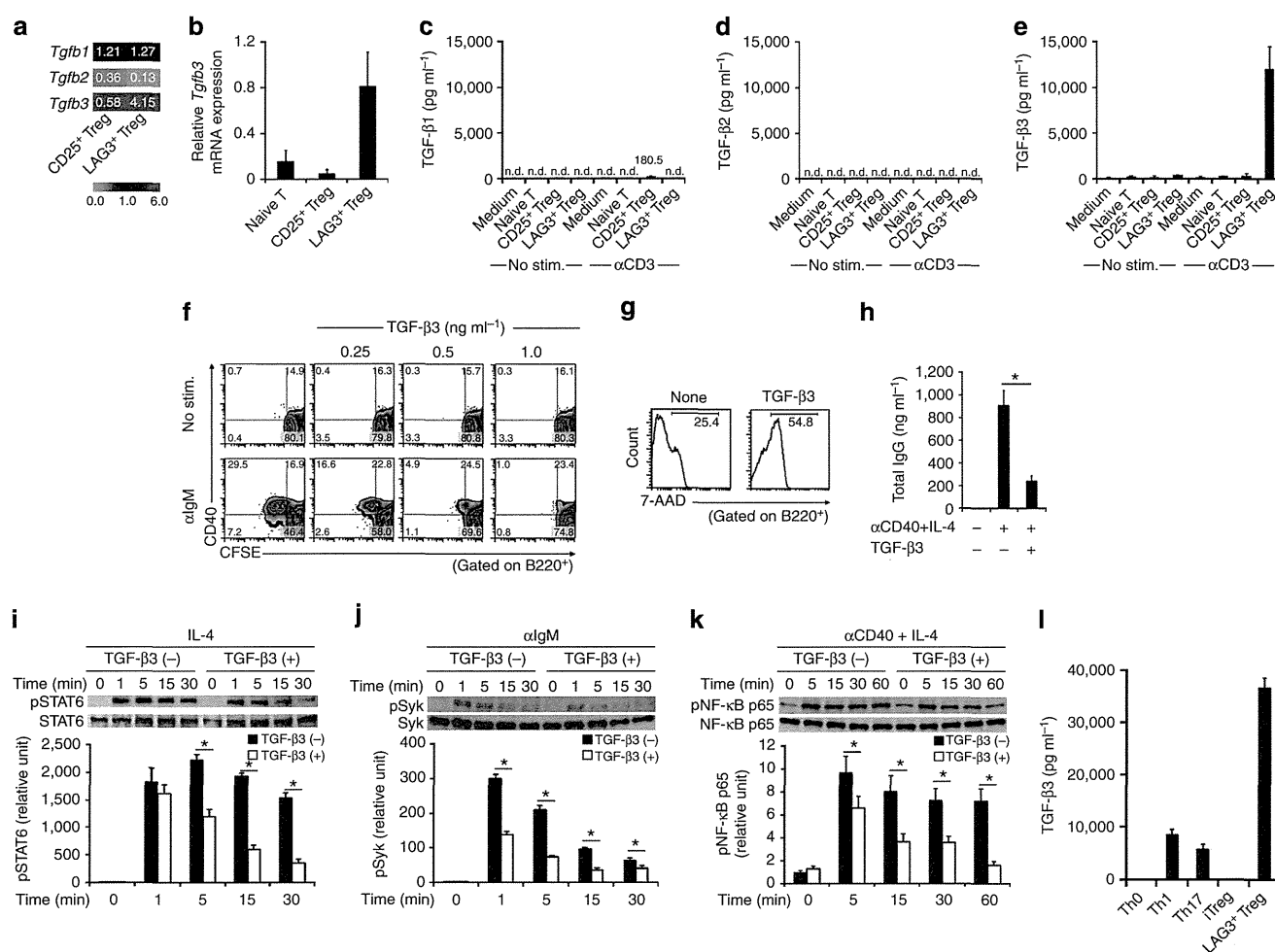


Figure 3 | LAG3⁺ Treg suppress B-cell activation through TGF- β 3. (a) Microarray comparisons of the gene expression profiles between B6 CD25⁺ Treg and B6 LAG3⁺ Treg. Normalized expression values from B6 CD4⁺ CD25⁻ CD45RB^{high} naive T cells are depicted according to the colour scale shown. (b) *Tgfb3* mRNA expression in sorted T-cell subsets taken from the spleens of B6 mice ($n = 3$ per group). (c–e) TGF- β 1, 2 and 3 protein levels in the culture supernatants of the indicated T-cell subsets from B6 mice determined using ELISA. Cells were seeded at 1×10^5 cells per well ($n = 4$ per group). (f) CFSE-labelled B cells were stimulated with or without anti-IgM mAb in the presence or absence of rTGF- β 3 (1 ng ml^{-1}) was assessed by 7-AAD ($n = 3$ per group). (g) Viability of anti-IgM-stimulated B cells in the presence or absence of rTGF- β 3 ($n = 3$ per group). (h) The effects of TGF- β 3 on total IgG production in the culture supernatants of anti-CD40/IL-4-stimulated B cells, determined as in Fig. 1d ($n = 3$ per group). * $P < 0.05$ (unpaired two-tailed Student's *t*-test). (i–k) STAT6 (i), Syk (j) and NF- κ B p65 (k) phosphorylation in stimulated B cells with or without rTGF- β 3, calculated as the ratio of phosphorylated to total protein levels ($n = 3$ per group). See also Supplementary Fig. 10. * $P < 0.05$ (unpaired two-tailed Student's *t*-test). (l) TGF- β 3 protein levels in the culture supernatants of freshly isolated B6 LAG3⁺ Treg or, naive B6 CD4⁺ T cells cultured under Th0, Th1, Th2 or Th17 conditions determined using ELISA. Cells were seeded at 3×10^5 cells per well ($n = 3$ per group). The means \pm s.d. are indicated.

transcription (STAT) 6, Syk and NF- κ B p65 in activated B cells (Fig. 3i–k and Supplementary Fig. 10). IL-4 produced by Th cells enhances the proliferation and survival of B cells while promoting immunoglobulin secretion and isotype switching via the activation of STAT6 (ref. 21). Activation of the tyrosine kinase Syk is critical for the cell signalling in response to B-cell receptor stimulation²². Activation of CD40, which is required for specific antibody production by antigen-stimulated B cells, induces phosphorylation of NF- κ B p65 (ref. 23). Therefore, TGF- β 3 inhibits several important pathways for B-cell functions. TGF- β 3 production is not limited to LAG3⁺ Treg because TGF- β 3 is also produced by developing Th17 cells in an IL-23-dependent manner²⁴. We found that Th1 cells produced TGF- β 3 in

addition to Th17 cells (Fig. 3l). However, LAG3⁺ Treg produced significantly greater amounts of TGF- β 3 than Th1 and Th17 cells.

Treatment with a TGF- β 3 blocking antibody cancelled the LAG3⁺ Treg-mediated suppression of antibody production and the development of T_{FH} and GCB in Rag1KO mice transferred with WT B cells and WT OT-II Th cells (Fig. 4a,b). Although TGF- β 1-LAP was detected on stimulated LAG3⁺ Treg (Supplementary Fig. 11a), TGF- β 1 blockade did not affect the suppressive activity of LAG3⁺ Treg for T_{FH} and GCB development and antibody production (Supplementary Fig. 11b,c). There was a possibility that reduction of antigen-specific IgG class antibody by LAG3⁺ Treg was related to the

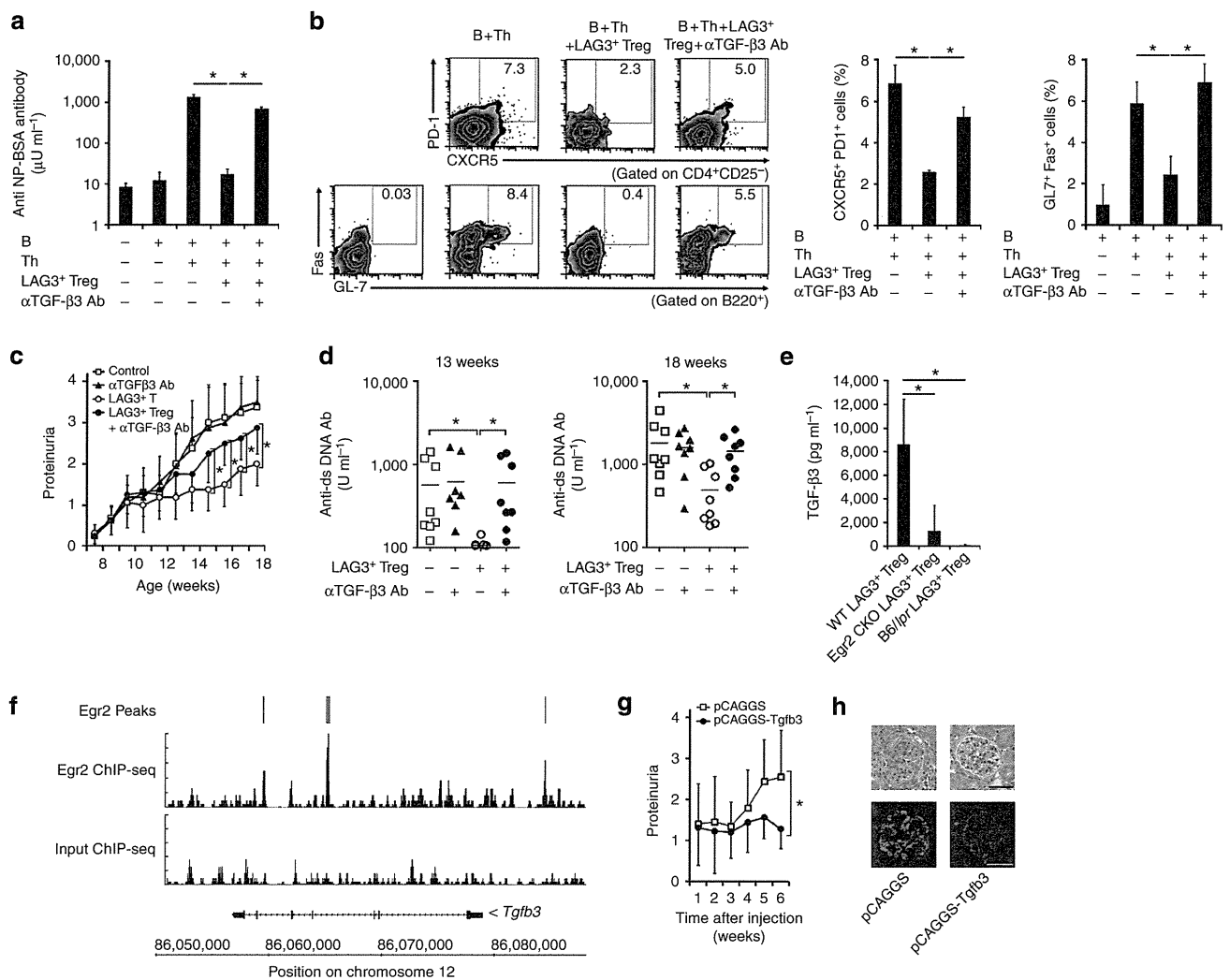


Figure 4 | TGF- β 3 ameliorates lupus manifestations. (a) *In vivo* blockade of LAG3⁺ Treg-mediated B-cell suppression by weekly injections of anti-TGF- β 3 blocking mAb (100 μ g per mouse) in NP-OVA-immunized Rag1KO mice transferred with B cells and T cells, as outlined in Fig. 1e ($n = 6$ per group). * $P < 0.05$ (Bonferroni post-test). (b) Flow cytometry plots and quantification of splenic CD4⁺CD25⁻CXCR5⁺PD-1⁺ T_{FH} and B220⁺GL-7⁺Fas⁺ GCB from the same mice as in a. * $P < 0.05$ (Bonferroni post-test). (c,d) Proteinuria progression (c) and serum levels of anti-dsDNA antibody (d) in MRL/+LAG3⁺ Treg-transferred MRL/lpr mice with or without a weekly injection of anti-TGF- β 3 mAb (100 μ g per mouse; $n = 8$ mice per group). Statistical significances in c were analysed with Mann-Whitney *U*-test, and d were analysed by Bonferroni post-test (* $P < 0.05$). (e) Production of TGF- β 3 by anti-CD3-stimulated LAG3⁺ Treg from WT, Egr2 CKO or B6/lpr mice, as in Fig. 3e ($n = 6$ per group). * $P < 0.05$ (Bonferroni post-test). (f) Distribution of Egr2-binding sites revealed by ChIP-seq in LAG3⁺ Treg. Egr2 ChIP-seq signal and Input ChIP-seq signal tracks are shown together with Egr2 peak calls (Egr2 Peaks). Peaks were identified using the Bioconductor package BayesPeak. (g) Proteinuria progression in MRL/lpr mice after i.v. injection with pCAGGS control ($n = 8$) or pCAGGS-Tgfb3 plasmid vector ($n = 7$). * $P < 0.05$ (Mann-Whitney *U*-test). (h) Representative images of kidney sections subjected to H&E staining (upper panels) and IgG immunofluorescent staining (lower panels) from the same mice as in Fig. 2c. Scale bars, 50 μ m. The experiments in e were repeated three times.

class switching to IgA because TGF- β and TGF- β receptor II (TGF β RII) signalling induce IgA class switching^{25–27}. However, no induction of antigen-specific IgA class antibody was observed by the co-transfer of WT LAG3⁺ Treg to Rag1KO mice transferred with WT B cells and WT OT-II Th cells (Supplementary Fig. 11d). TGF- β 3 blockade also abrogated the therapeutic effects of MRL/+ LAG3⁺ Treg in MRL/lpr mice (Fig. 4c,d), indicating a critical role for TGF- β 3. Intriguingly, TGF- β 3 production by Egr2-deficient LAG3⁺ Treg and Fas-mutated B6/lpr LAG3⁺ Treg was markedly reduced (Fig. 4e). Therefore, Egr2 and Fas are required for the production of TGF- β 3 and the B-cell-suppressive activity of LAG3⁺ Treg. Intriguingly, Fas-deficient B6/lpr mice contained LAG3⁺ Treg that express *Tgfb3* mRNA (Supplementary Fig. 12a,b), which indicated that Fas expression is not necessary for *Tgfb3* mRNA transcription in LAG3⁺ Treg. To further investigate the dependence of Egr2 on the transcription of *Tgfb3*, we performed Egr2 chromatin immunoprecipitation (ChIP)-seq analysis in LAG3⁺ Treg. The ChIP-seq analysis identified the binding sites of Egr2 in *Tgfb3* loci, suggesting direct regulation of the *Tgfb3* gene by Egr2 (Fig. 4f). The importance of TGF- β 3 for the control of lupus in MRL/lpr mice has also been verified by the observation that gene delivery of a TGF- β 3-expressing plasmid significantly improved proteinuria progression and renal pathology (Fig. 4g,h).

have been associated with SLE susceptibility¹. PD-1 provides negative co-stimulatory signals to both T cells and B cells^{28,29}, and PD-1-deficient (PD-1KO) mice develop a lupus-like disease³⁰. Furthermore, PD-1 and LAG3 synergistically regulate autoimmunity and tumour immunity^{31,32}. To examine the potential cooperation between TGF- β 3 and PD-1, we added TGF- β 3 to anti-IgM-stimulated B cells from B6, PD-1KO, Fas-deficient B6/lpr and FasL-deficient B6/gld mice. PD-1KO B cells, but not B6/lpr or B6/gld B cells, showed a partial resistance to the TGF- β 3-induced inhibition of cell division (Fig. 5a). Expressions of TGF- β RII, a receptor for TGF- β 3, on both T cells and B cells were not different among B6, PD-1 KO, B6/lpr and B6/gld mice (Supplementary Fig. 13). In accordance with a previous report that phosphorylated STAT6 is known to induce the expression of anti-apoptotic Bcl-xL in B-cell lines³³, TGF- β 3 suppressed the expression of Bcl-xL and Bcl-2a1 in activated WT B cells, but not in PD-1KO B cells (Fig. 5b). The cooperative suppression of B cells by TGF- β 3 and PD-1 underscores the importance of PD-1 expression on B cells for LAG3⁺ Treg-mediated suppression. The addition of an anti-PD-L1 blocking antibody reversed the LAG3⁺ Treg-mediated suppression of antibody production *in vitro* (Fig. 5c). Anti-NP antibody production was not suppressed by the co-transfer of WT LAG3⁺ Treg in Rag1KO mice transferred with PD-1KO B and WT OT-II Th cells; however, Rag1KO mice transferred with WT B cells and PD-1KO OT-II Th cells were susceptible to LAG3⁺ Treg-mediated suppression (Fig. 5d). Notably, TGF- β 3 significantly enhanced the expression of PD-1 on B cells stimulated with anti-IgM and anti-CD40 antibody (Fig. 5e). The interaction of PD-1 and its

LAG3⁺ Treg-mediated B-cell suppression requires PD-1. Genetic variants of the programmed cell death-1 (PD-1) gene

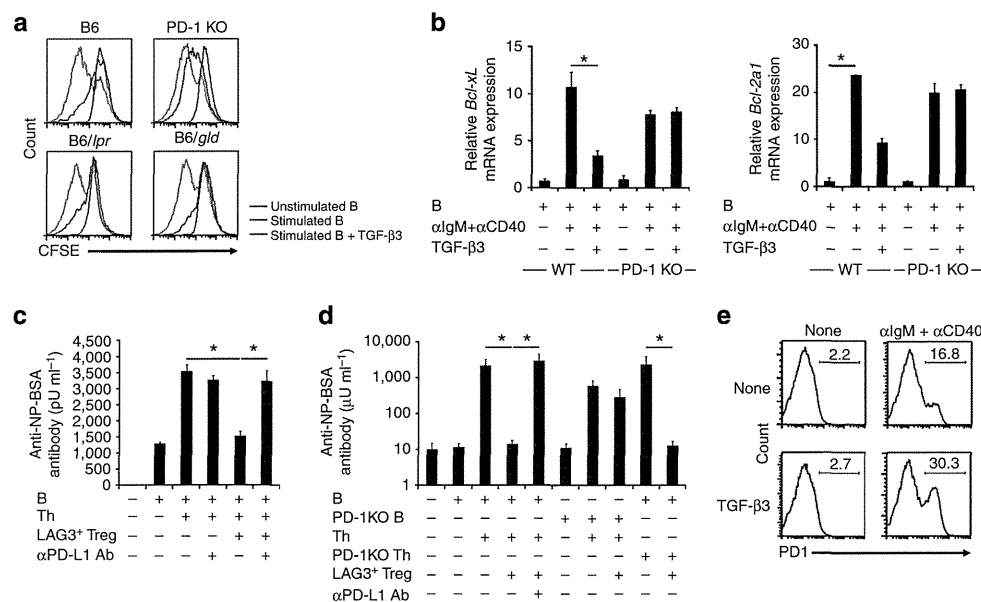


Figure 5 | PD-1 expression on B cells is important for the suppressive activity of LAG3⁺ Treg. (a) Resistance to TGF- β 3-mediated B-cell suppression in PD-1-deficient (PD-1KO) mice. CFSE-labelled B cells from B6, PD-1KO, B6/lpr or B6/gld mice were stimulated *in vitro* for 72 h with anti-IgM and anti-CD40 in the presence or absence of rTGF- β 3 (1 ng ml⁻¹). Histograms are gated on B220⁺ B cells. (b) *Bcl-xL* and *Bcl-2a1* mRNA levels in anti-IgM-stimulated B cells from WT or PD-1KO mice in the presence or absence of rTGF- β 3 (1 ng ml⁻¹; n = 3 per group). *P < 0.05 (unpaired two-tailed Student's t-test). (c) Blockade of LAG3⁺ Treg-mediated suppression of *in vitro* NP-specific antibody responses by anti-PD-L1 blocking mAb. B cells and Th cells purified from NP-OVA/alum-pre-immunized B6 mice and OT-II mice, respectively, were incubated with or without LAG3⁺ Treg from non-immunized OT-II mice in the presence or absence of anti-PD-L1 blocking mAb, and supernatants were analysed for anti-NP-BSA antibodies by ELISA (n = 6 per group). *P < 0.05 (Bonferroni post-test). (d) NP-specific antibody responses of Rag1KO mice injected with B6 B cells and OT-II Th cells from B6 or PD-1KO mice with or without LAG3⁺ Treg from B6 mice. Anti-PD-L1 blocking antibody (200 μ g per mouse) was injected *i.v.* every 3 days. Anti-NP-BSA antibody levels were determined as in Fig. 1e (n = 6 per group). *P < 0.05 (Bonferroni post-test). (e) PD-1 expression on B cells. B cells from B6 mice were stimulated *in vitro* for 72 h with or without anti-IgM and anti-CD40 antibodies in the presence or absence of rTGF- β 3 (1 ng ml⁻¹). Histograms are gated on B220⁺ B cells. Data are representative of three independent experiments. The means \pm s.d. are indicated.

ligand, PD-L1, may participate in the amplification of signalling cascade driven by TGF- β 3, which is essential for LAG3⁺ Treg-mediated B-cell suppression. These results confirm that PD-1 expression on B cells is required for LAG3⁺ Treg-mediated B-cell suppression. CD4⁺CD25⁻Egr2⁺T cells co-expressed both PD-L1, the ligand for PD-1, and LAG3 (Supplementary Fig. 14a,b). However, because PD-L1 and PD-L2 are expressed on various cell types including GCB³⁴, PD-L1 on LAG3⁺ Treg may not be the only source of ligands for PD-1 on B cells.

IL-27 induces TGF- β 3-producing Egr2⁺ T cells. IL-27 is a member of the IL-12/IL-23 heterodimeric family of cytokines produced by antigen-presenting cells (APCs). IL-27 has been identified as a differentiation factor for IL-10-producing Tr1 cells³⁵. We have previously reported that IL-27 induces Egr2 expression in CD4⁺ T cells, and Egr2 is required for the Blimp-1-mediated IL-10 production¹⁹. IL-27 treatment induced not only Egr2 but also a significant amount of TGF- β 3 protein (Fig. 6a,b). Egr2-deficient CD4⁺ T cells exhibited a substantial reduction in IL-27-induced production of TGF- β 3 (Fig. 6c), confirming the importance of Egr2 for the induction of TGF- β 3. The activation of specific STAT proteins in CD4⁺ T cells is associated with the differentiation of helper T-cell lineages. Although IL-27-mediated IL-10 induction requires both STAT1 and STAT3 (ref. 35), we previously found that IL-27-mediated induction of Egr2 is dependent on STAT3 (ref. 19). IL-27-mediated TGF- β 3

induction was impaired in STAT3-, not STAT1-, deficient CD4⁺ T cells (Fig. 6c), demonstrating similarity between Egr2 and TGF- β 3 in STAT3 dependency. Moreover, IL-27-treated CD4⁺ T cells significantly suppressed B-cell antibody production (Fig. 6d). In contrast, Egr2-deficient CD4⁺ T cells treated with IL-27 failed to suppress antibody production by B cells. IL-27-treated CD4⁺ T cells also exhibit suppressive activity for T_{FH} and GCB development and antibody production in a TGF- β 3-dependent manner (Fig. 6e,f). These results suggested that IL-27 induces CD4⁺ T cells that share several characteristics with LAG3⁺ Treg. Thus, IL-27-induced TGF- β 3-producing cells exhibit suppressive activity on humoral immunity in an Egr2-dependent manner.

Human LAG3⁺ Treg suppress antibody production. We identified CD4⁺CD25⁻CD45RA⁻LAG3⁺ T cells in CD4⁺ T cells from peripheral blood mononuclear cells (PBMCs) of healthy donors (Fig. 7a). Similar to murine LAG3⁺ Treg, human CD4⁺CD25⁻CD45RA⁻LAG3⁺ T cells expressed *EGR2*, *IL10* and *IFNG* (Fig. 7b) and produced significant amounts of IL-10 in response to TCR stimulation (Fig. 7c). The regulatory activity of human CD4⁺CD25⁻CD45RA⁻LAG3⁺ T cells was confirmed by the observation that they more efficiently suppressed antibody production when co-cultured with B cells and T_{FH} cells compared with CD4⁺CD25⁺CD127^{low}CD45RA⁻activated Treg³⁶ (Fig. 7d). Human CD4⁺CD25⁻CD45RA⁻LAG3⁺

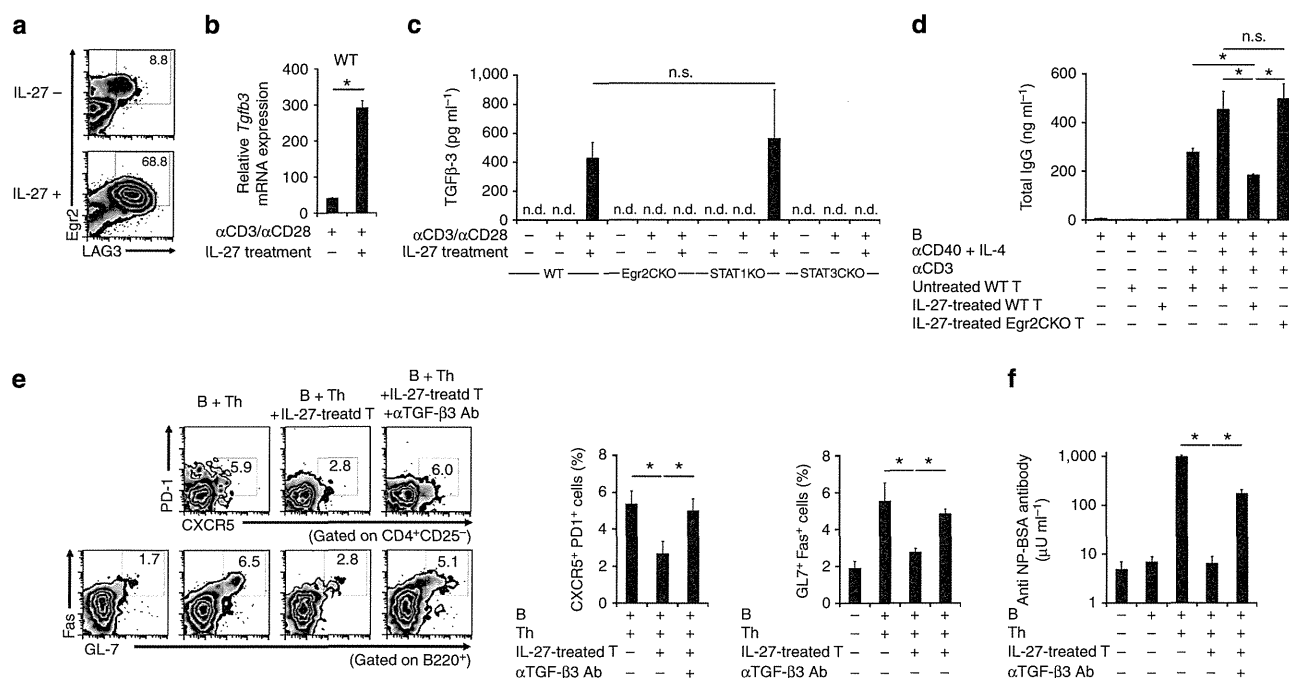


Figure 6 | IL-27 induces TGF- β 3-producing Egr2⁺ Treg from naive T cells. (a) IL-27-mediated induction of Egr2 and LAG-3 on CD4⁺ T cells. Freshly isolated naive CD4⁺ T cells were stimulated with anti-CD3/CD28 mAb in the presence or absence of IL-27. Cells were stained for Egr2 and LAG-3 expression on day 5. (b) Quantitative RT-PCR analysis of *Tgfb3* mRNA expression in naive WT CD4⁺ T cells activated as in a, assessed on day 3. **P* < 0.05 (unpaired two-tailed Student's *t*-test). (c) ELISA for TGF- β 3 in culture supernatants of activated naive WT, Egr2 CKO, STAT1 KO, or *Stat3*^{fl/fl} CD4-Cre⁺ (STAT3 CKO) CD4⁺ T cells as in a, assessed on day 5. **P* < 0.05 (Bonferroni post-test). (d) *In vitro* B-cell suppression by IL-27-treated naive WT CD4⁺ T cells. IL-27-treated or untreated naive WT or Egr2 CKO CD4⁺ T cells stimulated with anti-CD3 mAb were co-cultured with anti-CD40/IL-4-stimulated B cells. Total IgG was determined in culture supernatants on day 7 by ELISA (*n* = 3 per group). **P* < 0.05 (Bonferroni post-test). (e) Flow cytometry plots and quantification of splenic CD4⁺CD25⁻CXCR5⁺PD-1⁺ T_{FH} and B220⁺GL7⁺Fas⁺ GCB. B6 B cells and OT-II Th cells were transferred with or without IL-27-treated T cells into Rag1KO mice immunized twice with NP-OVA/alum, as in Fig. 1e (*n* = 6 per group). Anti-TGF- β 3 blocking antibody (100 μ g per mouse) was injected i.v. weekly. Numbers indicate the percentage of cells contained within the rectangular regions. **P* < 0.05 (Bonferroni post-test). (f) Serum levels of anti-NP-specific IgG1 antibody from the same mice as in e. **P* < 0.05 (Bonferroni post-test). n.d., not detected; n.s., not significant. Data are representative of three independent experiments. The means \pm s.d. are indicated.

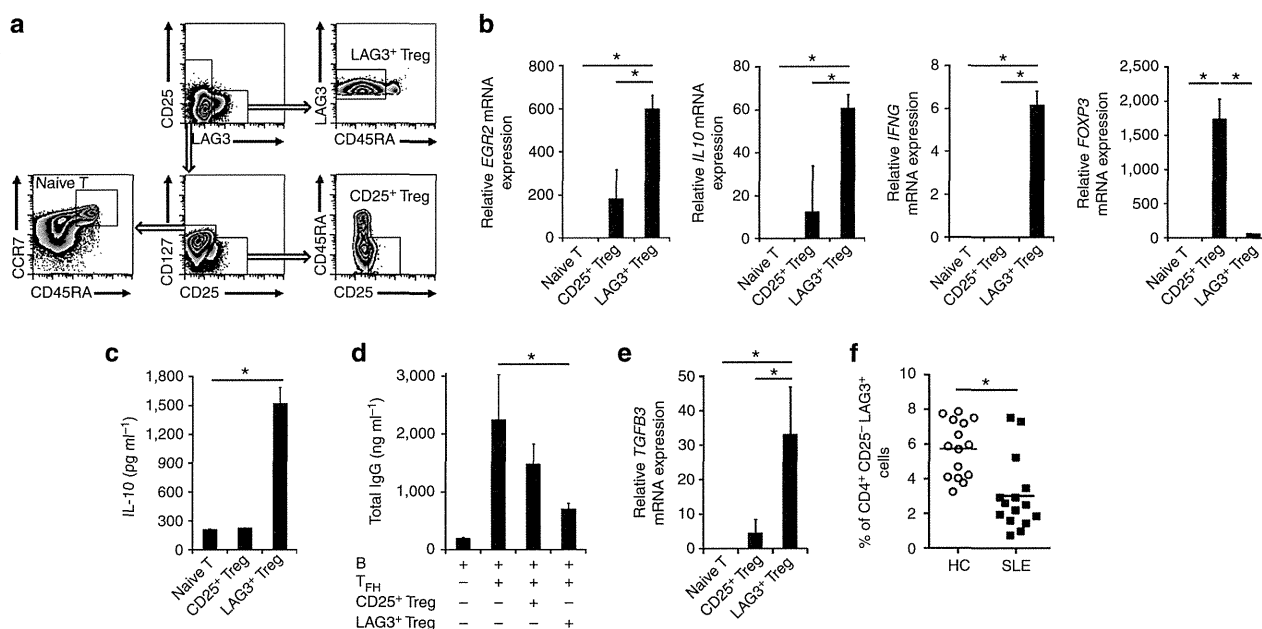


Figure 7 | Human CD4⁺ CD25⁻ CD45RA⁻ LAG3⁺ T cells suppress antibody production. (a) Gating strategy for CD4⁺ CD25⁻ CD127^{high} CCR7⁺ T cells (naive T), CD4⁺ CD25^{high} CD127^{low} CD45RA⁻ T cells (CD25⁺ Treg) and CD4⁺ CD25⁻ CD45RA⁻ LAG3⁺ T cells (LAG3⁺ Treg). Freshly isolated human PBMCs from healthy controls (HCs) were stained for CD4, CD25, CD45RA, CD127, CCR7 and LAG3, and the percentages of cells in each quadrant are indicated. (b) Quantitative RT-PCR analysis of *EGR2*, *IL10*, *IFNG* and *FOXP3* mRNA expression in anti-CD3 mAb-stimulated conditions for each CD4⁺ T-cell subset from HC ($n = 3$ per group). * $P < 0.05$ (Bonferroni post-test). (c) IL-10 protein levels in the culture supernatants on day 7 of the indicated T-cell subsets determined by ELISA ($n = 3$ per group). * $P < 0.05$ (unpaired two-tailed Student's t -test). (d) *In vitro* B-cell suppression by each CD4⁺ T-cell subset from HC. Each CD4⁺ T-cell subset was co-cultured with T_{FH} and staphylococcal enterotoxin B (SEB)-stimulated B cells. Total IgG was determined in culture supernatants by ELISA ($n = 3$ per group). * $P < 0.05$ (Bonferroni post-test). (e) *TGFβ3* mRNA expression in sorted T-cell subsets taken from HC ($n = 3$). * $P < 0.05$ (Bonferroni post-test). (f) Percentages of CD4⁺ CD25⁻ CD45RA⁻ LAG3⁺ T cells in each HC ($n = 15$) and SLE patients ($n = 15$) as in a. * $P < 0.05$ (unpaired two-tailed Student's t -test). The means \pm s.d. are indicated.

T cells expressed high levels of TGF- β 3 (Fig. 7e), suggesting that they suppress B cells through a mechanism identical to murine LAG3⁺ Treg. Therefore, we consider the CD4⁺ CD25⁻ CD45RA⁻ LAG3⁺ T-cell population as the human counterpart to murine LAG3⁺ Treg. We next assessed whether the frequency of LAG3⁺ Treg might be reduced in human systemic autoimmunity. The percentages of circulating LAG3⁺ Treg were significantly lower in the peripheral blood of SLE patients compared with healthy donors (Fig. 7f). These findings suggest that LAG3 expression could be used for tracking of the T-cell population with antibody-suppressing capacity in SLE patients.

Discussion

The results of the present study demonstrated that LAG3⁺ Treg suppress the development of GCB and T_{FH}, antibody production and disease progression in lupus-prone MRL/*lpr* mice. TGF- β 3, which is produced by LAG3⁺ Treg in Egr2- and Fas-dependent manners, plays a critical role in suppressing humoral immunity. As LAG3⁺ Treg also produce much higher levels of IL-10 than CD25⁺ Treg¹⁴, LAG3⁺ Treg are potent producers of regulatory cytokines. The pro-inflammatory role of TGF- β 3 was previously demonstrated by the observation that TGF- β 3 efficiently induces pathogenic Th17 cells^{24,37}. Our results have revealed a previously unrecognized role for TGF- β 3 in the control of autoimmunity. TGF- β 1 also exerts both pro-inflammatory and anti-inflammatory effects^{27,38,39}. In particular, TGF- β 1 induces B-cell apoptosis and reduces immunoglobulin production from activated human tonsil B cells^{40,41}. CD25⁺ Treg and Th3 regulatory cells^{42,43} are potent sources of TGF- β 1, and CD25⁺

Treg have been shown to suppress B-cell immunoglobulin synthesis through TGF- β 1 (ref. 44). However, the amount of TGF- β 1 produced by CD4⁺ T cells including CD25⁺ Treg is relatively limited (Fig. 3c), and it has been difficult to define the sources of TGF- β 1 that are relevant to immune suppression. Although it was demonstrated in a number of systems that TGF- β 1 and TGF- β 3 display clear isoform-specific biology, TGF- β 1 and TGF- β 3 showed comparable suppressive activity on B-cell responses (Supplementary Fig. 9a,b). In terms of helper T-cell development, TGF- β 3 is autonomously produced by Th17 cells during the development of pathogenic Th17 cells²⁴. In our setting, not only Th17 cells but also Th1 cells produced significant amounts of TGF- β 3; however, LAG3⁺ Treg produced greater amounts of TGF- β 3 compared with Th1 and Th17 cells (Fig. 3l). Therefore, the large amount of TGF- β 3 produced by LAG3⁺ Treg plays a significant role in the generation and maintenance of immune tolerance. As IL-10 strongly suppresses Th17 development and function⁴⁵, IL-10 produced by LAG3⁺ Treg may counteract the pro-inflammatory aspect of TGF- β 3.

We identified two molecules, Fas and Egr2, that are required for TGF- β 3 secretion in LAG3⁺ Treg. Egr2 deficiency in T cells and B cells results in a lupus-like syndrome, and Egr2 directly activates p21^{cip1} expression in CD44^{high} T cells and is involved in the control of Th1 and Th17 differentiation¹¹. The fact that Egr2 blocks the function of BATF, an AP-1 inhibitor required for the differentiation of Th17 cells, indicates that Egr2 is an intrinsic regulator of effector T cells¹². We showed here that Egr2 in T cells is important for the control of the development of T_{FH} and GCB (Fig. 1). Extrinsic functions of Egr2 for regulating humoral immunity were confirmed by our observation that transfer of

Egr2-sufficient LAG3⁺ Treg suppressed the excessive expansion of T_{FH} and GCB as well as antibody production in Egr2 CKO mice (Fig. 1). Fas controls T- and B-cell expansion by triggering apoptosis. The *lpr* gene, a mutation in the *Fas* gene in which the insertion of an early transposon results in a splicing error, interferes with peripheral T-cell tolerance, prevents Fas-dependent elimination of anergic B cells by CD4⁺ T cells and facilitates provision of T-cell help to autoreactive B cells. The impaired production of TGF-β3 by LAG3⁺ Treg may contribute to autoimmunity in MRL/*lpr* mice, as exogenous supplementation of TGF-β3 ameliorated the disease (Fig. 4g,h). These findings may indicate that Fas exhibits its tolerogenic activity via more complex regulatory pathways than previously thought.

IL-27, a differentiation factor for IL-10-producing Tr1 cells³⁵, induces CD4⁺Egr2⁺LAG3⁺ T cells¹⁹. However, the role of Tr1 cells in the regulation of B cells is not clear because CD46-induced Tr1 cells are more potent at enhancing immunoglobulin production compared with conventional T cells⁴⁶. In LAG3⁺Treg, IL-10 did not directly contribute to the control of B-cell response (Supplementary Fig. 7). Nevertheless, the linkage between IL-27 and the control of antibody production was suggested by the observation that overexpression of the IL-27 receptor, WSX-1, protects MRL/*lpr* mice from the development of autoimmune disease⁴⁷. It is notable that, whereas STAT1 and STAT3 are required for the induction of IL-10 by IL-27, the ability of IL-27 to promote Egr2 and TGF-β3 is STAT3-dependent (Fig. 6c). As STAT3-activating IL-6 also induces TGF-β3 production²⁴, STAT3 may play key roles for TGF-β3 induction in CD4⁺ T cells.

We here described potential cooperation of four molecules, Egr2, Fas, TGF-β3 and PD-1, in the control of humoral immunity. Activities of TGF-β family molecules are regulated by not only transcription but also protein-processing with proteases and integrins³⁷. Although the precise relationships remain to be clarified, Egr2 and Fas may regulate transcription and protein-processing of TGF-β3. The recruitment of SHP-2 phosphatase to the phosphorylated Tyr residue in PD-1 is in part responsible for the inhibitory effect through the dephosphorylation of signalling molecules belonging to the TCR or B-cell receptor pathways⁴⁸. Likewise, PD-1 signalling may modulate intracellular cascade downstream of TGF-β3.

Mouse and human CD25⁺ Treg share most of their features, although several differences exist regarding subset specificities and marker molecules. Human LAG3⁺ Treg express many key molecules in common with mouse LAG3⁺ Treg, and both suppress antibody production *in vitro*. Despite considerable efforts to clarify the contribution of CD25⁺ Treg to the development of SLE in patients, their role in generating disease remains elusive. However, Miyara *et al.*³⁶ reported that the number of CD4⁺CD25⁺CD127^{low}CD45RA⁻ activated Treg is decreased with a notable concomitant increase in the Foxp3^{low}CD45RA⁻ memory/effector-like non-Treg subset in active SLE. Several reports have described quantitative and qualitative reduction of CD25⁺ Treg in SLE⁴⁹. The functional impairment of immune-regulatory mechanisms may be crucial for the initiation and perpetuation of autoimmune disease, and a reduced frequency of LAG3⁺ Treg in SLE could also play a role in the dysregulated autoantibody production. Collectively, LAG3⁺ Treg may also be a regulatory mechanism for pathogenic autoantibody production in addition to CD25⁺ Treg. A deeper understanding of LAG3⁺ Treg will be useful for the treatment of autoantibody-mediated autoimmune diseases including SLE.

Methods

Mice. C57BL/6 (B6), C57BL/6-Fas^{pr^llpr} (B6/*lpr*), C57BL/6-Fas^{gld/gld} (B6/*gld*), MRL-Fas^{pr^llpr} (MRL/*lpr*) and MRL-Fas^{+/+} (MRL/+) mice were purchased from

Japan SLC. B6 recombinase-activating gene (*Rag*)-1-deficient (Rag1KO) mice, floxed-*Prdm1* (*Prdm1^{fl/fl}*) mice, *Il-10*-deficient (IL-10KO) mice, TCR transgenic OT-II mice (specific for the chicken ovalbumin peptide (amino-acid residues 323–339) in the context of MHC class II I-A^b) and TEa mice (specific for the Eα peptide (amino-acid residues 52–68) from the MHC class II I-Eα molecule in the context of I-A^b) were purchased from Jackson Laboratories. Floxed-*Stat3* (*Stat3^{fl/fl}*) mice were purchased from Oriental Bio Service (Japan). Rag1KO mice were housed in microisolator cages with sterile filtered air. B6-*Pdcd1*-deficient (PD-1KO) mice⁴⁷ were purchased from RIKEN BRC (Japan). Floxed *Egr2* (*Egr2^{fl/fl}*) mice were provided by Patrick Charnay (INSERM, France)⁴⁸. *Egr2* CKO mice (*Egr2^{fl/fl}* CD4-*Cre*⁺), *Prdm1* CKO mice (*Prdm1^{fl/fl}* CD4-*Cre*⁺) and STAT3 CKO (*Stat3^{fl/fl}* CD4-*Cre*⁺) mice were generated by crossing *Egr2^{fl/fl}* mice, *Prdm1^{fl/fl}* mice or *Stat3^{fl/fl}* mice with CD4-*Cre* transgenic mice on a B6 background, respectively. CD4-*Cre* transgenic mice (line 4196), originally generated by C. B. Wilson and colleagues, and *Stat1*-deficient (STAT1KO) mice were purchased from Taconic. Age- and sex-matched mice that were ≥ 7 weeks of age were used for all experiments. All animal experiments were approved by the ethics committee of the University of Tokyo Institutional Animal Care and Use Committee.

Reagents, antibodies and medium. The following reagents were purchased from BD Pharmingen; purified monoclonal antibody (mAb) for CD3ε (145-2C11), FasL blocking (MFL3), anti-CD40 (3/23), Fc block (anti-CD16/CD32, 1.5:100 dilution), fluorescein isothiocyanate (FITC) anti-CD45RB (16A, 1:100), FITC anti-Fas (Jo2, 2:100), phycoerythrin (PE) anti-CD45RB (16A, 1:100), APC-Cy7 anti-CD45RB (16A, 2:100), PE anti-LAG3 (C9B7W, 3:100), APC anti-LAG3 (C9B7W, 3:100), FITC anti-IgG1 (A85-1, 2:100), APC anti-IgG1 (A85-1, 2:100), FITC anti-GL7 (Ly-77, 0.5:100), FITC anti-CD25 (PC61, 2:100), PE anti-CD25 (PC61, 1:100), APC anti-CD25 (PC61, 1:100), APC-Cy7 anti-CD25 (PC61, 1:100), APC anti-CD4 (L3T4, 1:100), APC-Cy7 anti-CD4 (L3T4, 1:100), PE anti-CXCR5 (2G8, 5:100), APC-Cy7 anti-B220 (RA3-6B2, 1:100), PE anti-CD40 (3/23, 1:100), PE anti-PD-1 (J43, 1:100), biotinylated mAb for CD8a (53-6.7), CD19 (1D3), CD11c (HL3), CD45RB (16A), CD25 (7D4) and CXCR5 (2G8), streptavidin (SA)-FITC antibody (Ab), SA-APC and SA-APC-Cy7. Alexa Fluor 488 anti-LAG3 mAb (C9B7W, 1:100) and FITC anti-PD-L1 mAb (MIH6, 2:100) were purchased from AbD Serotec. Qdot605 anti-CD4 mAb (RM4-5, 2:100) and SA-Qdot605 were purchased from Invitrogen. PE anti-PD-L1 mAb (MIH5, 1:100), PE anti-Egr2 mAb (erongr2, 1:100), PE anti-PD-L1 mAb (10F.9G2, 1:100) and APC anti-B220 mAb (RA3-6B2, 1:100) were purchased from eBioscience. NP(13)-OVA and NP(9)-BSA were purchased from Biosearch Technologies. SA-conjugated microbeads were purchased from Miltenyi Biotec. FITC anti-mouse IgG Ab was purchased from Sigma. BrilliantViolet421 anti-B220 mAb, APC anti-mouse LAP (Tgf-β1, 2:100) and anti-PD-L1 blocking mAb (10F.9G2) were purchased from Biologend. Alexa 488 Fluor anti-GFP mAb was purchased from Medical & Biological Laboratories. Recombinant TGF-β1 (rTGF-β1) and three were purchased from Miltenyi Biotec, and rTGF-β2, anti-TGF-β1 blocking polyclonal Ab (MAB240), anti-TGF-β3 blocking polyclonal Ab (MAB234), APC anti-mouse TGF-βRII (1.5:100), APC goat IgG and rIL-27 were purchased from R&D Systems.

For human studies the following anti-human mAbs were used: V450 anti-hCD4 (RPA-T4, 2:100), V500 anti-hCD4 (RPA-T4, 2:100; both from BD Biosciences), PerCP-Cy5.5 anti-human CD3 (UCHT1, 2:100), Brilliant Violet 421 anti-hCD25 (BC96, 3:100), APC anti-hCD19 (HIB19, 2:100), PerCP-Cy5.5 anti-hCD4 (OKT4, 5:100), Alexa Fluor 647 anti-hCD197 (G043H7, 3:100), APC-Cy7 anti-human CD45RA (HI100, 3:100; all from BioLegend), Alexa Fluor 488 anti-hCD25 (BC96, 4:100) and PE-Cy7 anti-hCD127 (eBioRDR5, 1:100; all from eBioscience). PE anti-hLAG3 polyclonal Ab (10:100) was purchased from R&D Systems.

T cells and B cells were cultured in RPMI-1640 medium supplemented with 10% FBS, 100 μg ml⁻¹ L-glutamine, 100 U ml⁻¹ penicillin, 100 μg ml⁻¹ streptomycin and 50 μM 2-mercaptoethanol (all purchased from Sigma).

Generation of Egr2-GFP mice. The bacterial artificial chromosome (BAC) clone RP23-88D4, which contained the entire genomic *Egr2* locus, was obtained from BC Libraries (Invitrogen). This clone was modified for the insertion of an enhanced GFP gene (eGFP) with an SV40 polyA sequence at the initiation codon in Egr2 exon 1 using the Red/ET recombination system. The Egr2-eGFP construct linearized with *P*I-SceI was injected into the pronuclei of fertilized zygotes from B6 mice and transferred to pseudopregnant females. Offsprings were screened for genomic integration by PCR of tail DNA using the following Egr2 promoter primer: forward 5'-AGACCGCATTACTCTTATCACCAG-3', SV40polyA-specific primer: reverse 5'-TGAGTTTGACAAACCACTAG-3' (PCR product size: 2.1 kb). Mice were generated by breeding F1 heterozygous transgenic males with WT females.

Cell purification. Briefly, spleens were cut into pieces and digested with collagenase type IV (Sigma-Aldrich). Red blood cells were lysed with hypotonic shock induced by brief exposure to ammonium chloride with potassium lysis buffer, followed by immediate isotonic restoration. Surface staining was performed in ice-cold PBS with 2% fetal calf serum in the presence of an FcR-blocking antibody (anti-mouse CD16/CD32 mAb). To obtain highly purified CD4⁺ T cells, single-cell suspensions were first purified by negative selection with

magnetic-activated cell sorting (MACS; Miltenyi Biotec) using anti-B220 mAb, anti-CD19 mAb, anti-CD11c mAb and anti-CD8a mAb. To obtain highly purified CD4⁺CD25⁻CD45RB^{low}LAG3⁺ T cells, CD45RB^{high} cells were subsequently depleted with anti-CD45RB mAb. After FcR-blocking, the prepared cells were stained with mAbs specific for CD4, CD25, CD45RB and LAG3 in order to isolate CD4⁺CD25⁻CD45RB^{low}LAG3⁺ T cells (LAG3⁺ Treg), CD4⁺CD25⁺ T cells (CD25⁺ Treg), CD4⁺CD25⁻LAG3⁻ Th cells or CD4⁺CD25⁻CD62L^{hi}CD44^{low} (naive T). Cells for intracellular anti-Egr2 staining were stained using the Foxp3 Staining Buffer Set (eBioscience) according to the manufacturer's protocol. The purities of MACS- and FACS (FACSVantage SE (Becton-Dickinson) or MoFlo XDP (Beckman Coulter))-sorted cells were >90% and >99%, respectively.

Immunization. NP-OVA in alum was prepared by mixing NP(13)-OVA (Biosearch Technologies) in PBS with alum (Pierce) at a 1:1 ratio for 30 min at 4 °C. The immunization with NP-OVA/alum was performed by intraperitoneal injection.

Adoptive transfer of LAG3⁺ Treg from WT mice into Egr2 CKO mice.

FACS-purified 2 × 10⁵ LAG3⁺ Treg from B6 mice were injected intravenous (i.v.) into 10-week-old Egr2 CKO mice pre-immunized with or without 100 µg NP-OVA/alum 1 day before the cell transfer. The development of T_{FH} and GCB in the spleen was analysed with FACS 7 days after the cell transfer. Serum levels of anti-BSA-NP antibodies were analysed using enzyme-linked immunosorbent assay (ELISA) 7 days after the immunization, as described above.

Generation of mixed-BM chimeras. BM cells were harvested by flushing the femurs and tibias of donor mice with RPMI medium. Lethally irradiated Thy1.2⁺ Egr2 CKO mice (700 rad) were reconstituted with 2 × 10⁶ Thy1.2⁺ Egr2 CKO BM cells or a mixture of 1 × 10⁶ Thy1.2⁺ Egr2 CKO and 1 × 10⁶ Thy1.1⁺ WT BM cells. The recipient mice were used in the analyses 6 weeks after BM transfer.

Localization of splenic Egr2-GFP⁺ T cells. Spleens from Egr2-GFP mice and Foxp3-GFP mice were rapidly frozen in Tissue-Tec OCT compound (Sakura Finetek, Japan) with liquid nitrogen, and then were cut into 5-µm sections with a cryostat microtome. Sections were fixed for 10 min in 4% paraformaldehyde (Wako) and were preincubated in PBS with 2% BSA and 0.1% saponin, and then incubated with the following primary antibodies for 30 min: PE anti-mouse CD4 (5:160), Brilliant Violet 421 anti-mouse B220 (2:160), FITC anti-GFP (3:160). Images were acquired with a fluorescence microscope (Olympus BX53). The frequencies of Egr2-GFP⁺CD4⁺ T cells and Foxp3-GFP⁺CD4⁺ T cells were evaluated by counting the numbers per field; each field was 0.01 mm².

B-cell isolation and proliferation. Splenic B cells were purified by negative selection with MACS using a B-cell isolation kit (Miltenyi Biotec) according to the manufacturer's protocol. The purity of MACS-sorted B cells was >95% positive for B220 staining. B cells were labelled with 5 µM 5-(and 6-) carboxyfluorescein diacetate succinimidyl ester (CFSE; Dojindo) at 37 °C for 10 min, and then were stimulated with 10 µg ml⁻¹ anti-IgM F(ab)₂ (Jackson ImmunoResearch Laboratories) for 72 h with or without rTGF-β1, 2 or 3 for B-cell proliferation assays. Cells were stained with anti-B220 mAb, 7-Amino-Actinomycin D (7-AAD; Biolegend, 3:100) and PE anti-CD40 mAb. The percentages of viable 7-AAD-negative CFSE-diluted B220⁺CD40⁺ B cells and dead 7-AAD-positive B220⁺ cells were assessed using FACS.

In vitro B-cell activation co-culture assays. The wells of 96-well flat-bottomed plates were coated with 10 µg ml⁻¹ anti-CD3 mAb in 100 µl per well of PBS and incubated overnight at 4 °C. The wells were washed, and MACS-purified B cells with or without each FACS-purified T-cell subset (LAG3⁺ Treg, CD25⁺ Treg or CD4⁺CD25⁻CD44^{lo}CD62L^{hi} naive T cells) or IL-27-treated CD4⁺ T cells described below were plated immediately into the coated wells at a density of 1 × 10⁵ cells per well for each cell type in RPMI medium as described above alone or with 10 µg ml⁻¹ anti-CD40 mAb (3/23) + 10 µg ml⁻¹ rIL-4 (Cell Signaling Technology) supplemented with or without rTGF-β3 (1 ng ml⁻¹). B cells undergoing apoptosis on day 3 and total IgG production in the culture supernatants on day 7 were determined using the Annexin V Apoptosis Detection Kit (BD Pharmingen) and a mouse IgG ELISA Quantitation Set (Bethyl Laboratories), respectively, according to the manufacturer's protocol.

Adoptive transfer studies in Rag1KO mice and TEa mice. MACS-purified 2 × 10⁵ B cells from B6 or PD-1KO mice and FACS-purified 2 × 10⁵ Th cells from OT-II or PD-1KO OT-II mice were injected i.v. into Rag1KO mice in combination with or without FACS-purified 1 × 10⁵ LAG3⁺ Treg from B6, Egr2 CKO, B6/lpr, B6/gld or IL-10KO mice, or 3 × 10⁵ IL-27-treated T cells. Control mice received PBS. Mice were subsequently immunized with 100 µg NP-OVA/alum 24 h after the cell transfer. Mice were re-immunized with 50 µg NP-OVA/alum 14 days after the first immunization. Where indicated, the day after the cell transfer, mice were injected i.v. with an anti-FasL blocking antibody (200 µg per mouse), anti-TGF-β1

blocking antibody (100 µg per mouse) or anti-TGF-β3 blocking antibody (100 µg per mouse) at weekly intervals, or an anti-PD-L1 blocking antibody (200 µg per mouse) every 3 days. Serum anti-NP antibody levels were analysed with ELISA, and splenocytes were analysed with FACS 7 days after the re-immunization. To examine the *in vivo* suppressive activity of LAG3⁺ Treg in non-lymphopenic conditions, TEa mice were employed. TEa TCR is supposed to have low cross-reactivity²⁷ and TEa mice possess very few CD25⁺ Treg⁵⁰. FACS-purified 2 × 10⁵ Th cells from OT-II mice were injected i.v. into TEa mice in combination with or without FACS-purified 1 × 10⁵ LAG3⁺ Treg from B6 mice. Control mice received PBS. Mice were subsequently immunized with 100 µg NP-OVA/alum 24 h after the cell transfer. Serum levels of anti-BSA-NP antibodies were analysed with ELISA 14 days after the immunization, as described above.

Quantification of NP-specific antibody responses. Anti-NP IgG antibody levels were quantified by ELISA using NP(9)-BSA (Biosearch Technologies) as the capture antigen in the *in vitro* or *in vivo* antibody production assay, respectively. ELISA plates were prepared using the Immuno-Tek ELISA construction system (Zepto Matrix) according to the manufacturer's protocol. Following the incubation with sample serum or media, the plates were developed with horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG1, or HRP-conjugated goat anti-mouse IgA (SouthernBiotech), and TMB substrate. Serially diluted pooled sera from NP(13)-OVA-immunized B6 mice were included as controls on each plate. The concentrations of the anti-NP IgG1 antibody were estimated by comparisons with standard curves constructed from pooled sera.

Adoptive transfer studies in MRL/lpr mice. Eight-week-old MRL/lpr mice were randomly assigned to specific treatment groups. Ten-week-old MRL/lpr mice in the treatment group were injected i.v. with LAG3⁺ Treg, LAG3⁻ T cells, CD25⁺ Treg or naive T cells (1 × 10⁵ cells each) obtained from MRL/+ mice. The three-time injection group (MRL/+ LAG3⁺ Treg x3, MRL/+ CD25⁺ Treg x3 and MRL/lpr LAG3⁺ Treg x3) was adoptively transferred i.v. with 1 × 10⁵ LAG3⁺ Treg at weekly intervals (10, 11 and 12 or 13, 14 and 15 weeks of age, respectively); the mice were 10 weeks of age at the time of the first injection). Each T-cell subset was first enriched by MACS and then sorted by FACS (on the basis of the expression of CD4, CD25, CD45RB and LAG3) as described above. Mice in the control group received PBS. Where indicated, the day after the first cell transfer, mice were injected i.v. with an anti-TGF-β3 blocking antibody (100 µg per mouse) at weekly intervals. Mice were killed at 18 weeks of age to examine pathological alterations. Anti-ds DNA antibodies were measured using a mouse anti-ds DNA ELISA Kit (Shibayagi) at 13 and 18 weeks of age, according to the manufacturer's protocol.

Urinary protein analysis. Proteinuria was assessed semiquantitatively using dip sticks (Albustix, Bayer) at weekly intervals (0 = none; 1 = 30–100 mg dl⁻¹; 2 = 100–300 mg dl⁻¹; 3 = 300–1,000 mg dl⁻¹; 4 ≥ 1,000 mg dl⁻¹).

Histological analysis. MRL/lpr mice were killed at 18 weeks of age. Renal pathology was graded by standard methods for glomerular inflammation, proliferation, crescent formation and necrosis as described elsewhere⁴⁹. The proportion of glomeruli was evaluated by examining at least 80 glomeruli per section by an examiner blind to the experimental conditions. Interstitial and tubular changes were also noted. Scores from 0 to 4 (where 0 represents 'no damage' and 4 represents 'severe') were assigned for each of these features.

In vitro NP-specific antibody responses. B6 and OT-II mice were immunized with 100 µg NP-OVA/alum. Mice were re-immunized with 100 µg NP-OVA/alum 3 weeks after the first immunization. MACS-purified 2 × 10⁵ B cells from pre-immunized B6 mice and FACS-purified 1 × 10⁵ CD4⁺CD25⁻LAG3⁻ Th cells from pre-immunized OT-II mice were seeded in round-bottom 96-well plates 7 days after the re-immunization in combination with or without FACS-purified 1 × 10⁵ LAG3⁺ Treg from non-immunized OT-II mice in the presence or absence of 10 µg ml⁻¹ anti-PD-L1 blocking mAb (10F.9G2) or 10 µg ml⁻¹ anti-FasL blocking mAb (MFL3). The culture supernatants were harvested at 3 weeks, and anti-NP antibody levels were analysed with ELISA, as described above.

RNA isolation, cDNA synthesis and quantitative real-time PCR. Total T-cell RNA was prepared using an RNeasy Micro Kit (Qiagen). RNA was reverse-transcribed to cDNA with random primers (Invitrogen) and SuperScript III in accordance with the manufacturer's protocol (Invitrogen). To determine the cellular expression level of each gene, quantitative real-time PCR analysis was performed using an iCycler (Bio-Rad). The PCR mixture consisted of 25 µl of SYBR Green Master Mix (Qiagen), 15 pmol of forward and reverse primers and the cDNA samples in a total volume of 50 µl (ref. 14). Relative RNA abundance was determined on the basis of control mouse β-actin or human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) abundance.

DNA microarray analysis. Total RNA of CD4⁺CD25⁺, CD4⁺CD25⁻CD45RB^{low}LAG3⁺ and CD4⁺CD25⁻CD45RB^{high}LAG3⁻ FACS-purified T cells