Research Article

BACH2 in TRegs Limits the Number of Adipose Tissue Regulatory T Cells and Restrains Type 2 Immunity to Fungal Allergens

Amanda Contreras,1 Darin L. Wiesner,2 Brock Kingstad-Bakke,1 Woojong Lee,1 John P. Svaren,3 Bruce S. Klein,2 and M. Suresh1

1Department of Pathobiological Sciences, University of Wisconsin-Madison, Madison, 53706 WI, USA
2Department of Medical Microbiology and Immunology, University of Wisconsin-Madison, Madison, 53706 WI, USA
3Department of Comparative Biosciences, University of Wisconsin-Madison, Madison, 53706 WI, USA

Correspondence should be addressed to M. Suresh; sureshm@vetmed.wisc.edu

Received 21 June 2022; Accepted 19 July 2022; Published 5 August 2022

Academic Editor: Mitesh Dwivedi

Copyright © 2022 Amanda Contreras et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

FoxP3+ regulatory T cells (Tregs) are essential for self-tolerance and moderating tissue-damaging inflammation. Tregs that develop and mature in the thymus are classified as central Tregs or effector Tregs based on whether Tregs predominantly inhabit secondary lymphoid organs (central Tregs) or tissues (effector Tregs). By generating mice that are conditionally deficient for Bach2 in peripheral Tregs, we have examined the role of Bach2 in regulating Treg homeostasis and effector functions. Unlike global and T cell-specific Bach2-deficient mice, Treg-specific Bach2 ablation did not result in unprovoked TH2 inflammation in the lungs. However, Bach2 deficiency in Tregs led to augmented expressions of IRF4, BATF, and GATA3 and a significant increase in the accumulation of ST2 (IL-33R)+ve effector Tregs in the spleen and visceral adipose tissue (VAT) but not in the lungs. Enhanced Bach2-deficient Treg numbers in VAT was not linked to hyperresponsiveness to exogenous IL-33 in vivo. Most strikingly, Treg-specific Bach2 deficiency resulted in enhanced fungal protease-induced Type 2 allergic inflammation in the lungs, with no detectable effect on Type 1 responses to systemic or respiratory viral infections. In summary, we ascribe vital roles for Bach2 in peripheral Tregs: as a transcriptional checkpoint to limit precocious differentiation into effector Tregs in lymphoid tissues and as a regulator of the functional program that restrains Type 2 but not Type 1 inflammation in lungs. Results presented in this manuscript implicate dysregulated Tregs in the pathogenesis of airway hypersensitivities, asthma, and other allergic disorders.

1. Introduction

Foxp3+ regulatory T (Treg) cells are a vital T-cell subset that enforce self-tolerance and mitigate inflammatory pathology during immune responses to foreign antigens [1]. Peripheral Treg cells are divided into two distinct subtypes based on their anatomical distribution and function: central (cTregs) and effector Tregs (eTregs) [2, 3]. The majority of Treg cells in the secondary lymphoid tissues are cTregs that express high levels of lymphoid homing receptors CD62L and CCR7 and have trafficking patterns similar to naïve conventional T cells. By contrast, eTregs constitute a small fraction of Tregs in the lymphoid tissues, but the majority of Tregs in the peripheral tissues. Unlike cTregs, the eTregs display an activated effector phenotype (e.g., low expression levels of CD62L and CCR7, but high levels of CD44, inducible T cell costimulatory [ICOS], glucocorticoid-induced tumor necrosis factor receptor [GITR], and KLRG-1) with trafficking patterns similar to conventional effector T cells (i.e., traffic between blood and peripheral non-lymphoid tissues) [4, 5]. These Foxp3+ eTregs have been further classified into T_{H1}-Tregs, T_{H2}-Tregs, and T_{H17}-Tregs depending upon their polarity and co-opted expression of T-bet, GATA-3, and STAT-3 transcription factors, respectively [6]. This pattern of polarized differentiation endows each eTreg subset with distinct migratory and functional properties tailored to specifically balance inflammatory
responses driven by T_{h1}, T_{h2}, or T_{h17} effector T cells. While IL-2R signaling is clearly necessary for eTreg cell differentiation [7–9], activation and polarization of cTregs into eTreg cell subsets are driven by TCR signaling and the inflammatory milieu in the peripheral tissues [10–13].

Tissue Tregs are a specialized subset of effector Tregs that reside in non-lymphoid tissues [14] and display characteristics including expression of transcription factors and homing or effector molecules that allow them to specialize for their specific tissue environment [14]. The IL-33 receptor ST2 thus far appears to be a universal feature of tissue Tregs that distinguishes them from classical lymphoid-organ Tregs [14]. While it is known that BATF and IRF4 are required for the induction of key components of the effector Treg transcriptional program, it has recently been demonstrated that BATF and IRF4 are required for the induction of eTregs or tissue Tregs from cTregs are not well understood. Bach2 is a transcription repressor that plays an integral role in the maintenance of T cell quiescence, differentiation, and generation of memory T cells [17–19]. Using global Bach2-deficient mice, we and others have shown that Bach2 plays crucial roles in the differentiation and competitive fitness of Tregs and protects against fatal T_{h12}-driven eosinophilic crystalline pneumonia [20, 21]. Further, it has been reported that ablation of Bach2 in all T cells leads to dysregulated T_{h2} immunity and pulmonary inflammation in mice [22–24]. Until recently, Treg-specific role of Bach2 in maintaining the homeostasis of peripheral Tregs or suppressing T_{h1}/T_{h2} immunity was unknown. Two reports have since shown that Bach2 restrains the differentiation of eTregs by repressing genes regulated by transcription factors such as IRF-4 [25–27]. Here, we report that ablation of Bach2 in peripheral Tregs leads to enhanced differentiation of effector/tissue Tregs in the lymphoid tissues, increased numbers of visceral adipose tissue (VAT) Tregs, and increased activation of T cells in response to normal homeostatic cues. Further, we show that loss of Bach2 in Tregs does not affect the development of T_{h1}/T_{h2} cells during mucosal or systemic viral infections but instead augments the differentiation of IL-5 and IL-13-producing T_{h2} cells during fungal allergen-induced inflammation in the lungs. These findings provide new insights into the Treg-specific roles for Bach2 in (i) regulating the homeostasis of splenic and VAT eTregs and (ii) restraining fungal protease-induced Type 2 immunity but not virus-induced Type 1 immunity.

2. Materials and Methods

2.1. Mice. Mice with a FoxP3+ Treg-specific deletion of Bach2 were generated by breeding floxed Bach2 mice [19] with B6.129(Cg)-Foxp3^{flh/wVFFV/piere2Abf/J} mice [28] to create Bach2^{loxp/loxp}FoxP3-Cre mice. Either littermate WT or C57BL/6J mice were used as controls. The mice used in these studies were housed in specific-pathogen-free conditions in University of Wisconsin-Madison animal facilities. All animal experiments were conducted in accordance with approved protocols of the institutional animal care committee.

2.2. Viral Infections and Protease Treatment. The Armstrong strain of lymphocytic choriomeningitis virus (LCMV) was administered intraperitoneally at 2×10^6 plaque-forming units (PFU) per mouse to six- to eight-week-old mice; virus-specific T cell responses in spleen were quantified at day 8 after LCMV infection. The influenza virus strain A/PR/8/34 H1N1 (PR8) was administered intranasally at 100 PFU per mouse. Mice were euthanized, and lungs were harvested 10 days after PR8 infection. For fungal protease treatment, 10-week-old mice were anesthetized to receive intratracheal doses of 25 μg of fungal protease from Aspergillus fumigatus (Sigma #P4032) on days 1, 2, 7, and 14. On day 15 after the initial treatment, mice were injected with 3 μg of anti-CD45 antibody intravenously immediately prior to euthanasia and lung harvest.

2.3. IL-33 In Vivo Administration. Mice received intraperitoneal injections of 0.5 μg of recombinant murine IL-33 (Peprotech) or PBS on days 0, 2, 4, and 6 [29]. On the 8th day of IL-33 treatment, mice were euthanized to harvest lungs, VAT, and spleens.

2.4. IL-33 In Vitro Treatment. Spleens from naïve Bach2^{loxp/loxp}FoxP3-Cre and WT mice were processed into single cell suspensions and were plated at a concentration of 5 × 10^5 cells/well in a 96-well plate. Prior to plating cells, wells were coated with 1 μg/mL anti-CD3. Cells were cultured for 72 hours with IL-2 (10 ng/mL) (BD Biosciences) and anti-CD28 (2 μg/mL) in either the presence or absence of 1 ng/mL IL-33. Cells were then collected and processed for flow cytometry.

2.5. Flow Cytometry. Single-cell suspensions of mononuclear cells from lung, VAT, and spleen were prepared using standard procedures. Lung and VAT were digested in 2 mg/mL Collagenase D (Sigma) for 30 minutes while rotating at 37°C. The digested tissues were then homogenized using the GentleMACS Dissociator (Miltenyi). Single-cell suspensions were first stained for viability with a LiveDead stain (eBioscience). Cells were then resuspended and stained with antibodies diluted in a staining buffer of either 2% BSA in PBS or in Brilliant Stain Buffer (BD Biosciences), depending on the combination of antibodies used. The fluorochrome-labeled antibodies that were used against cell-surface antigens include the following: CD8, CD4, CD25, CD44, CD62L, CD127, CD69, GITR, CTLA-4, KLRG-1, CCR7, CD103, CXCR3, ST2, CD27, CD11b, CD90.2, CD64, Siglec-F, PD-L1, CD19, CD11c, Ly6G, p80 TCR, and TCR-β. Fluorochrome-labeled antibodies that were used against intracellular antigens include the following: IFN-γ, IL-4, IL-5, IL-17A, BAF, IRF4, GATA3, Foxp3, Ki-67, Eomes, Tbet, and Helios. These antibodies were purchased from BD Biosciences (San Jose, CA), Biolegend (San Diego, CA), or eBioscience (San Diego, CA). Fluorochrome-conjugated tetrmers for LCMV epitopes (D\textsuperscript{3}/NP396, D\textsuperscript{3}/GP33, I-A\textsuperscript{b}/GP66) and for PR8/H1N1 epitopes (D\textsuperscript{3}/NP366, D\textsuperscript{3}/PA224, I-A\textsuperscript{b}/NP311) were provided by the NIH Tetramer Core Facility (Emory University, Atlanta, GA). Samples were acquired with a BD LSRFortessa (BD Biosciences), and
2.6. Intracellular Cytokine Staining. To induce cytokine production in cells for intracellular cytokine staining, cells were stimulated directly ex vivo with human recombinant IL-2 (10 U/well) (BD Biosciences) and the epitope peptide at 0.1ug/mL. Depending on the experiment, cells were either stimulated with PMA/Ionomycin (Tonbo Biosciences) or LCMV peptides (NP396, GP33, and GP276) or PR8/H1N1 peptides (NP366, PA224, and NP311) (thinkpeptides, Prolimmune Ltd.). Cells undergo 5 hours of stimulation at 37°C in the presence of brefeldin A (1ug/mL, GolgiPlug, BD Biosciences). After stimulation, cells were stained for cell surface antigens, fixed, and permeabilized using the Cytotix/Cytoperm kit (BD Sciences). After fixation/permeabilization, cells were incubated with fluorochrome-labeled antibodies targeted against cytokines. For the fungal protease studies, cells extracted from the lungs were restimulated directly ex vivo with 100 ug/well of heat inactivated fungal protease for 6 hours. Brefeldin A was added for the last 4 hours of the restimulation. The cells were then fixed, permeabilized, and stained for intracellular cytokine expression as described above.

2.7. Intracellular Staining for Transcription Factors and Ki-67. To stain for intracellular proteins such as transcription factors and Ki-67, cells were fixed and permeabilized with the FoxP3 Staining Kit (eBioscience) using the manufacturer’s protocol. After fixation/permeabilization, cells were incubated with fluorochrome-labeled antibodies targeted against transcription factors and Ki-67. After staining, cells were analyzed with a BD LSRFortessa flow cytometer.

2.8. Statistical Analyses. Data statistics were calculated with Prism software (GraphPad Software, La Jolla, California, USA). Student’s two-tailed t-test and one-way ANOVA analyses were used to calculate the statistical significance of differences between groups, and significance was defined at p < 0.05.

3. Results

3.1. Bach2 Restrains the Activation and Differentiation of Effector Tregs and Maintains Homeostasis of Naïve and Activated/Memory T Cells. We and others have previously reported that global Bach2 deficiency leads to aberrant differentiation of Tregs and activated/memory phenotype of peripheral T cells [20, 21]. To investigate the effect of conditional Bach2 deficiency in mature Tregs, we bred flexed Bach2 mice with Rudensky’s Foxp3-Cre mice to generate the Bach2loxp/loxpFoxP3-Cre mice. Loss of Bach2 in Tregs did not alter the frequency or total numbers of Foxp3+ve Tregs in spleen, which suggested that Bach2 is not required for the development and/or maintenance of these cells (Figures 1(a) and 1(b)). Next, we assessed whether Treg-specific Bach2 deficiency dysregulated the homeostasis of naïve and activated effector Tregs. Notably, Bach2-deficient Tregs exhibited enhanced expression of CD44, CD127, CD69, and GITR, which was strongly suggestive of skewed differentiation favoring an effector Treg phenotype (Figure 1(c)) [4, 5]. Clearly, there was a significant increase in the percentages and numbers of the activated/effector (CD44hi/CD62Llo/CCR2lo) Bach2-deficient Tregs, as compared to their WT counterparts (Figure 1(d)). Further, there were increased frequencies and total numbers of Bach2-deficient Tregs that expressed high levels of KLRG1, CD127, and CD69.

Next, we investigated whether loss of Bach2 in Tregs affected the homeostasis of classical naïve and activated/memory T cells in trans. We found an increase in the percentages of CD4 and CD8 T cells with an activated/effector phenotype (CD44hi/CD62Llo) in the spleens of Bach2loxp/loxpFoxP3-Cre mice compared to WT counterparts (Figure 1(e)). In addition, we observed a decrease in the percentages of CD4 T cells with a naïve phenotype (CD44lo/CD62Lhi) (Figure 1(e)). Next, we were interested in whether Bach2-deficient Tregs altered the production of cytokines from conventional T cells. Splenic CD8 and CD4 T cells had a substantial increase in the percentages and total numbers of cells that produced IFNγ in the Bach2loxp/loxpFoxP3-Cre mice compared to WT mice. Additionally, CD4 T cells from Bach2loxp/loxpFoxP3-Cre mice displayed higher frequencies of IL-13 and IL-17A-producing cells as well as total cells that produced IL-17A (Figures 1(f) and 1(g)). These augmented levels of cytokine-producing cells suggest that Bach2-deficient Tregs are unable to repress the activation and differentiation of Th1 (IFNγ), Th2 (IL-13), and Th17 (IL-17A) effector CD4 T cells. Taken together, these data suggested that Bach2 plays an essential role in maintaining quiescence/naïveté by restraining the differentiation of naïve Tregs into effector Tregs. Furthermore, loss of Bach2 expression in Tregs led to the development of activated/memory T cells, which suggests that the ability of Tregs to limit activation of classical T cells requires Bach2 expression in Tregs.

3.2. Bach2 Represses Tissue Treg-like Cell Differentiation in Secondary Lymphoid Tissues. Apart from the well-defined subsets of naïve and effector Tregs in secondary lymphoid tissues, recent work has identified a distinct subset of Tregs termed as tissue Tregs that can be found in non-lymphoid tissues such as lungs, skeletal muscle, and lamina propria [14]. While these tissue Tregs display the prototypical effector surface phenotype markers such as CXCR3 and CD103 and express transcription factors BATF, IRF4, and GATA3, they possess certain properties that make each tissue Treg unique. However, tissue Tregs possess a distinguishing surface marker that has been found on all tissue-residing Tregs populations currently examined: ST2. ST2 is a receptor for IL-33 and is expressed on Tregs that preferentially accumulate in non-lymphoid tissues [30, 31]. It was of interest to determine whether Bach2 deficiency dysregulated the differentiation of tissue Tregs in lymphoid tissues. As expected, a modest fraction of Tregs expressed CD103, CXCR3, and ST2 in spleens of WT mice. Surprisingly, however, we found that Bach2 deficiency in Tregs led to a substantial increase in the frequencies of CD103hi, CXCR3hi, and ST2hi Tregs in spleen, as compared to their WT counterparts. The total numbers of CD103hi and ST2hi Tregs were significantly higher in the spleens of Treg-specific Bach2-deficient mice than in WT mice.
Figure 1: Continued.
Figure 1: Continued.

**Figure Legend:**
- **Panel (e):** Graph showing the percentage of CD4 and CD8 T cells stained for CD44 and CD62L in WT and Bach2^loxp/loxp^FoxP3-Cre mice.
- **Panel (f):** Graph showing the percentage of IFN-γ^+^ CD4 and CD8 T cells in WT and Bach2^loxp/loxp^FoxP3-Cre mice.

**Table:**
- |       | CD44 LO | CD44 HI | CD62L HI | CD62L LO |
  |-------|---------|---------|-----------|-----------|
  | **WT** | 64.7 ± 4.2 | 20.5 ± 2.5 | 48.0 ± 9.6 | 38.0 ± 2.9 |
  | **Bach2^loxp/loxp^FoxP3-Cre** | 48.0 ± 9.6 | 38.0 ± 2.9 | 7.6 ± 1.6 | 10.2 ± 2.5 |

**Notes:**
- IFN-γ^+^ CD4^+^ T cells:
  - WT: 5 ± 2%
  - Bach2^loxp/loxp^FoxP3-Cre: 10 ± 2%

- IFN-γ^+^ CD8^+^ T cells:
  - WT: 2 ± 1%
  - Bach2^loxp/loxp^FoxP3-Cre: 5 ± 2%
Histograms are gated on Foxp3 +ve Treg cells and show staining for the indicated markers in WT (shaded) and Bach2 loxp/loxp FoxP3-Cre mice. Indeed, we found statistically significant increases of BATF, IRF4, and GATA3 expression in Bach2-deficient mice (Figure 2). It is possible that the increased numbers of tissue Treg-like cells in the spleen of Treg-specific Bach2-deficient mice could result from defects in trafficking of these cells to non-lymphoid tissues. To address this possibility, we quantified Tregs in lungs and liver of Treg-specific Bach2-deficient mice. Data in Figure 2 show that the percentages and numbers of Tregs in lungs and liver (data not shown) of Treg-specific Bach2-deficient mice were comparable to those in WT mice. Therefore, it is likely that tissue Tregs are accumulating in spleen of Treg-specific Bach2-deficient mice due to increased differentiation and not driven by defective trafficking of these cells from spleen to non-lymphoid tissues.

We next investigated whether Bach2 deficiency-induced enhancement in the development of ST2+ve tissue Tregs was associated with altered expression of tissue Treg fate determining transcription factors BATF, IRF4, and GATA3. Indeed, we found statistically significant increases of BATF, IRF4, and GATA3 expression in Bach2-deficient Tregs compared to their WT counterparts only in spleen, but not in lungs (Figure 2(b)). Overall, our studies have revealed an unexpected increase in the ST2+CD103+CXCR3+ tissue Treg-like cells in the spleen of mice harboring a Treg-specific Bach2 deletion. These data suggest that Bach2 exerts a transcriptional block in the differentiation of tissue Treg-like cells in the lymphoid tissues.

3.3. Bach2 Limits the Numbers of Visceral Adipose Tissue Tregs. Tissue Tregs have been most thoroughly characterized in the VAT, and transcription factors BATF and IRF-4 bind to the ST2 promoter and promote ST2 expression [15]. Since Bach2-deficient Tregs express elevated levels of ST2, BATF, and IRF-4 (Figure 2), it was of interest to investigate whether Bach2 regulated the homeostasis of VAT Tregs. Analysis of Tregs in VAT showed that Bach2 deficiency led to a significant increase in the numbers of CD103+ve and CXCR3+ve Tregs in VAT (Figure 3(b)); the numbers of ST2+ve Tregs were not significantly altered in VAT of Treg-specific Bach2-deficient mice. Next, we examined whether Bach2 deficiency-induced increased numbers of VAT Tregs was associated with elevated expression of transcription factors that drive the VAT Treg program. We found that Bach2-deficient VAT Tregs exhibited significantly higher levels of GATA3 and IRF4 compared to WT Tregs (Figure 3(c)).

ST2 is the cellular receptor for IL-33, and it is known that ST2 and IL-33 are required for the development of VAT Tregs [15]. Mechanistically, IL-33 enhances ST2 expression by inducing GATA-3 phosphorylation and subsequent recruitment of GATA-3 to the ST2 locus. Because Bach2-deficient Tregs have higher expression of ST2 in the spleen and express elevated levels of GATA3 and IRF4, we explored whether Bach2-deficient Tregs are more poised to the effects of IL-33 and therefore more readily differentiate into tissue Tregs. As expected, in vitro exposure of WT splenic Tregs to IL-33 induced ST2 expression in a fraction of Tregs (Figure 3(d)). While the percentages of ST2-expressing Tregs were already higher among untreated Bach2-deficient Tregs, IL-33 treatment further increased the percentages of ST2+ve Tregs (Figure 3(d)). The magnitude of ST2 induction was comparable in WT and Bach2-deficient Tregs, but the IL-33-induced expressions of GATA3, IRF4, and BATF were greater in Bach2-deficient Tregs, as compared to those in WT Tregs (Figure 3(d)), which suggest that Bach2 restraints IL-33-induced expression of these transcription factors in Tregs.

**Figure 1:** Bach2 regulates classical and regulatory T cell homeostasis. Splenocytes from naïve WT and Bach2loxp/loxp FoxP3-Cre mice were stained for intracellular and extracellular antigens and analyzed by flow cytometry. (a) Frequency and (b) number of Foxp3+ve Treg cells. (c) Histograms are gated on Foxp3+ve Treg cells and show staining for the indicated markers in WT (shaded) and Bach2loxp/loxp FoxP3-Cre mice (line) mice. (d) Frequency and numbers of Tregs expressing the indicated molecules. (e) Activation of classical T cells was analyzed by staining with anti-CD44 and anti-CD62L; contour plots are gated on CD4 or CD8 T cells. (f), (g) Splenocytes were stimulated for 5 hrs in vitro with PMA and ionomycin in the presence of Brefeldin A, and cytokine production by CD4 and CD8 T cells was measured by intracellular staining for (f) IFN-γ and (g) IL-13 and IL-17A. This experiment was repeated three times with similar results, 4-5 mice per group. *p < 0.05.
Figure 2: Continued.
Exogenous IL-33 administration has been reported to drive the expansion of VAT Tregs in vivo [15]. Here, we investigated whether Tregs from WT or Bach2-deficient Treg mice respond differently to IL-33 treatment in vivo. IL-33 treatment significantly augmented the numbers of Tregs only in the VAT, but not in the spleen or lungs (Figure 3(e)) of both WT and Bach2lox/loxFoxP3-Cre mice. The IL-33-responsive population of Tregs included CD103⁺, CXCRC3⁺, and ST2⁺ Tregs in both groups of mice (Figure 3(f)). Mechanistically, IL-33-driven expansion of Tregs in VAT was associated with elevated percentages of proliferating Ki67⁺ Tregs (not shown) in both WT and Bach2lox/loxFoxP3-Cre mice. As compared to PBS-treated controls, the magnitude of increase in the number of Tregs was similar for WT and Bach2lox/loxFoxP3-Cre mice. Taken together, data in Figure 3 strongly suggest that Bach2 deficiency did not alter the IL-33-driven proliferative expansion of VAT Tregs.

3.4. Bach2-Deficiency in Tregs Does Not Alter the Development of Th1 or Th17 Cells during an Acute Viral Infection. Viral infections typically trigger Type 1 immune responses and Tregs play a crucial role in restraining the effector phase of T cell responses to mitigate inflammation and the associated tissue damage. Data in Figure 1 showed that conventional CD8 and CD4 T cells from Bach2lox/loxFoxP3-Cre mice displayed greater activated/effector phenotypic and functional properties, as compared to conventional T cells from WT mice. The elevated levels of IFNγ production in CD4 T cells of Bach2lox/loxFoxP3-Cre mice suggested that deficiency for Bach2 in Tregs might lead to deregulated Type 1 immunity. Therefore, it was of interest to determine whether (1) deregulated Treg homeostasis in Bach2lox/loxFoxP3-Cre mice is altered by viral infections and (2) Bach2 deficiency in Tregs affects the development of Th1 and Th17 type T cells during an acute viral infection. First, we infected cohorts of WT and Bach2lox/loxFoxP3-Cre mice with lymphocytic choriomeningitis virus (LCMV) and assessed Tregs in spleen at day 8 after infection. The percentages and total number of Foxp3⁺ Tregs in spleens of LCMV-infected Bach2lox/loxFoxP3-Cre mice were similar to those in WT mice (Figure 4(a)). Next, we evaluated the expression of surface markers associated with an activated/effector Treg phenotype in LCMV-infected WT and Bach2lox/loxFoxP3-Cre mice. There were no significant changes in the percentages of CD44, CD62L, CXCRC3, CD27, CD127, and KLRG-1-expressing Tregs in unaftered mice. Likewise, the numbers of Foxp3⁺ Tregs between the groups were similar (Figure 4(c)). Taken together, data in Figure 4 suggest that Treg-specific Bach2 deficiency did not alter the development of CD4 and CD8 T cell responses during an acute LCMV infection.

Next, we assessed the effect of Treg-specific Bach2 deficiency on CD4 and CD8 T cell responses to LCMV infection. The frequency and numbers of LCMV-specific CD8 T cells in Bach2lox/loxFoxP3-Cre mice were similar to those in WT mice. Likewise, the numbers of CD4 T cells specific to the LCMV GP66 epitope in WT mice were comparable to those in Bach2lox/loxFoxP3-Cre mice (Figure 4(c)). To investigate if Bach2 deficiency altered cytokine production by LCMV-specific CD4 and CD8 T cells, we measured antigen-induced IFN-γ production ex vivo. We did not find significant differences in the percentages of LCMV-specific IFN-γ-producing CD8 and CD4 T cells in spleens between WT and Bach2lox/loxFoxP3-Cre mice (Figure 4(d)). Taken together, data in Figure 4 suggest that Treg-specific Bach2 deficiency did not alter the development of CD4 and CD8 T cell responses during an acute LCMV infection.

Next, we assessed the effect of Treg-specific Bach2 deficiency on Th1/Th17 responses to a mucosal infection with influenza A virus (IAV). First, we infected Bach2lox/loxFoxP3-Cre and WT mice with PR8 strain of IAV and analyzed Tregs in the lung on day 10 post-infection. We did not find significant differences in the number of Tregs between the Bach2lox/loxFoxP3-Cre and WT mice in the lung (Figure 4(e)). Likewise, there were no changes in the percentages of CD44, CD62L, CXCRC3, CD103, CD27, CD69, CD127, and KLRG-1-expressing Tregs in the lung or BAL of influenza virus-infected Bach2lox/loxFoxP3-Cre mice (Figure 4(f)).

Next, we investigated whether Bach2-deficiency in regulatory T cells altered the activation and expansion of effector
Figure 3: Continued.
Figure 3: Bach2 limits the size of tissue regulatory T cell population in the VAT. Mononuclear cells isolated from the VAT of WT and Bach2^{loxp/loxp}-FoxP3-Cre mice were analyzed by flow cytometry. Tregs in the VAT of naïve WT and Bach2^{loxp/loxp}-FoxP3-Cre mice were analyzed for (a) their frequency and numbers, (b) tissue Treg marker expression, and (c) MFI levels for BATF, IRF4, and GATA3. (d) Splenocytes from naïve WT and Bach2^{loxp/loxp}-FoxP3-Cre mice were stimulated in vitro with anti-CD3, anti-CD28, and IL-2 in the presence of absence of IL-33 for 72 hrs. Bar graphs display frequencies of ST2^{+ve} and BATF^{+ve}, IRF4^{+ve}, or GATA3^{+ve} Tregs. (E-H) WT and Bach2^{loxp/loxp}-FoxP3-Cre mice received four injections of IL-33, and the indicated tissues were analyzed on day 8. (e) Frequency and number of Treg cells in the VAT, lungs, and spleen. (f) Number of CXCR3-, CD103-, and ST2-positive Tregs in the VAT. This experiment was repeated two times with similar results, 5 mice per group. *p < 0.05.
Figure 4: Continued.
Figure 4: Continued.
CD8 and CD4 T cells during an influenza infection. No significant differences in the frequencies or total number of influenza-specific CD8 or CD4 T cells were seen in the lung and BAL of Bach2<sup>loxp/loxp</sup>FoxP3-Cre mice (Figure 4(g)). To determine if there were any changes in the production of IFN-γ by influenza-specific CD4 and CD8 T cells, we stimulated lymphocytes in the lung with corresponding influenza peptides and measured IFN-γ production. Although stimulation with the PA224 peptide elicited greater percentages of IFN-γ-producing CD8 T cells in Bach2<sup>loxp/loxp</sup>FoxP3-Cre mice.
Cre mice (Figure 4(h)), Bach2-deficiency in Tregs had minimal effects on the activation or cytokine production of CD8 or CD4 T cells, during a mucosal T_{H1}/T_{C1} response to influenza virus.

3.5. Bach2 Deficiency in Tregs Exacerbates Fungal Protease-Induced T_{H2} Immunity. Genome-wide association studies have linked variations in Bach2 gene to asthma in humans, and global Bach2 deficiency in mice leads to unprovoked T_{H2} immunity and fatal eosinophilic crystalline pneumonia [20, 32, 33]. Additionally, Bach2 deficiency in all T cells leads to spontaneous development of T_{H2}-driven lung disease [24]. We observed that Treg-specific Bach2 deficiency did not result in overt lung pathology, which suggested that Bach2 expression in conventional T cells is sufficient to protect against spontaneous T_{H2} lung disease. However, it was unknown whether Bach2 deficiency in Tregs affected (1) their responses to allergic inflammation and (2) T_{H2} responses and susceptibility of mice to allergen-provoked inflammation in the lungs. To investigate the effect of Treg-specific Bach2 deficiency on T_{H2} immunity, we utilized a model of allergic inflammation in lungs of protease-treated Bach2 loxp/loxpFoxP3-Cre mice (Figure 5(e)). Taken together, data in Figure 5 strongly linked variations in Bach2 gene to asthma in humans, and we utilized a model of allergic inflammation in the lungs of protease-treated Bach2 loxp/loxpFoxP3-Cre mice to elicit a potent T_{H2} response in the lungs [34]. The immunological response of Bach2^{loxp/loxp}FoxP3-Cre and WT mice to Aspergillus protease administration is shown in Figure 5. As shown in Figure 5(a), protease administration induced substantial increases in the numbers of Tregs in the lungs of WT mice, and protease-induced Treg accumulation was markedly accentuated in lungs of Bach2^{loxp/loxp}FoxP3-Cre mice. The increased numbers of Tregs in lungs of protease-treated Bach2^{loxp/loxp}FoxP3-Cre mice included CXCR3^{+ve}, CD103^{+ve}, ST2^{+ve}, and KLRG-1^{+ve} Tregs (Figure 5(b)), and this was associated with elevated levels of IRF4 and GATA3 (Figure 5(c)). We also compared the accumulation of inflammatory cells in the lungs of protease-treated WT and Bach2^{loxp/loxp}FoxP3-Cre mice (Figure 5(d)). Overall, lungs of protease-treated Bach2^{loxp/loxp}FoxP3-Cre mice contained higher numbers of monocytes, monocyte-derived DCs, CD103^{+ve} DCs, inflammatory DCs, alveolar macrophages, neutrophils, and eosinophils, as compared to protease-treated WT mice. Next, we assessed whether Treg-specific Bach2 ablation affected the T_{H1}/T_{H2} polarization of protease-reactive CD4 T cells in the lungs. Significantly more IL-5- and IL-13-producing CD4 T cells were detected in the lungs of protease-treated Bach2^{loxp/loxp}FoxP3-Cre mice than in WT mice (Figure 5(e)). Taken together, data in Figure 5 strongly suggest that Bach2 plays a critical role in promoting the ability of Tregs to limit T_{H2} immunity and allergic inflammation in the lungs.

4. Discussion

It has been established that ST2^{+ve} tissue Tregs are a specialized subset of effector Tregs that reside primarily in non-lymphoid tissues [14, 15], but not in the spleen [15, 16, 35, 36]. In this study, we report a surprising finding that a substantive proportion of Bach2-deficient Tregs in spleen expressed conventional activation markers and elevated levels of tissue Treg markers CXCR3, CD103, and ST2. The increase in the numbers of effector/tissue Tregs in spleen of Treg-specific Bach2-deficient mice cannot be explained by defective homing to the peripheral tissues because lungs and VAT contained normal or greater numbers of Tregs. Apart from IL-2R and TCR signaling, development of eTregs is driven by inflammatory milieu in the peripheral tissues [7–13]. Governed by extracellular cues, transcription factors BATF and IRF4 maintain tissue Tregs, orchestrate the effector Treg transcriptional program, and promote expression of ST2 [15]. Subsequently, IL-33 induces GATA-3 phosphorylation, which binds to the ST2 locus to enhance ST2 gene expression [16]. The expression of ST2 in T_{H2} [37, 38] and tissue Tregs [16, 39] is reliant on a positive feedback loop where IL-33 induces GATA3 recruitment to the ST2 locus, IL1rl1. Studies by Vasanthakumar et al. demonstrated that IRF4 and BATF binding to the IL1rl1 loci is required for maintaining VAT Treg identity, and administration of IL-33 amplifies the number of Tregs in the VAT [15]. Our discovery of ST2^{+ve}GATA3^{+ve} Bach2-deficient Tregs in the spleen prompted us to further investigate whether Bach2-deficient splenic Tregs are hypersensitive to IL-33, as compared to their WT counterparts. We find that IL-33 exposure in vitro further increased the expression of ST2 in splenic Bach2-deficient Tregs. This increase is impressive considering that Bach2-deficient Tregs already express elevated basal levels of ST2 compared to WT Tregs. This increase of ST2 expression was associated with increased frequencies of GATA3-, BATF-, and IRF4-expressing cells, which suggest that Bach2 might repress GATA3/BATF/IRF4-driven expression of ST2 and subsequent differentiation of Tissue Tregs in the spleen. Furthermore, Bach2 restrains differentiation of eTregs by competing with, or directly repressing BATF, IRF4, and GATA3 expression in response to extracellular cues [21, 24–27], which in turn can limit ST2 expression and eTreg development.

Most notably, Treg phenotypes were largely normal in non-lymphoid tissues such as lungs, but Bach2-deficiency in Tregs resulted in substantive increase in the numbers of CXCR3^{+ve} and CD103^{+ve} Tregs in the VAT. Although there was not any detectable change of ST2 expression, Bach2-deficient Tregs exhibited higher levels of GATA3 and IRF4, overall suggesting that Bach2 is needed to repress tissue Treg differentiation in the VAT. In the present study, we found that in vivo IL-33 treatment increased the numbers of Tregs in VAT of Bach2^{loxp/loxp}FoxP3-Cre and WT mice and that these IL-33-responsive cells were CD103^{+ve}, CXCR3^{+ve}, and ST2^{+ve} Tregs. Notably, WT Treg levels of GATA3, BATF, and IRF4 were induced to levels that were comparable to those of IL-33-treated Bach2-deficient Tregs. In contrast to our in vitro experiment with splenic Tregs, IL-33 treatment did not induce significant in vivo changes in the splenic Treg population in Bach2-deficient mice. This difference could be attributed to the fact that Bach2-deficient Tregs already demonstrate a higher basal level of expression of tissue Treg phenotype compared to WT and, in vivo, it could take a higher concentration to exacerbate this phenotype than the regimen that we used. Alternatively, IL-33-induced Tregs in spleen might have already re-localized to VAT.

Despite displaying a robust effector phenotype, Bach2-deficient Tregs seemed to be incapable of restraining the
Figure 5: Continued.
activation of conventional T cells. Higher numbers of CD8 and CD4 T cells in Treg-specific Bach2-deficient mice exhibited an activated effector phenotype as determined by CD44 and CD62L expression. Further, these CD8 and CD4 T cells secreted substantial amounts of pro-inflammatory cytokines. Specifically, higher percentages of CD4 T cells from Treg-specific Bach2-deficient mice secrete IFNγ, IL-13, and IL-17A, suggesting that Tregs require Bach2 expression to repress the activation and differentiation of TH1, TH2, and T H17 cells, respectively. These findings led us to explore whether Bach2 deficiency in Tregs is unable to control Type 1 and Type 2 responses in vivo. Acute viral infections typically induce Type 1 responses, and it was of interest to assess whether Bach2 deficiency altered the effect of LCMV infection on Tregs and whether Treg-specific Bach2 deficiency led to enhanced T cell responses to an LCMV infection. Surprisingly, Treg-specific Bach2 deficiency has no detectable effect on the numbers of Tregs or the magnitude of virus-specific CD8/CD4 T cell response or the cytokine-producing ability of virus-specific CD8/CD4 T cells. Unlike in an LCMV infection, Treg-specific Bach2 deficiency limits the expansion of Tregs in mice infected with Plasmodium chabaudi [26]. An emerging theme from our studies is that loss of Bach2 expression in Tregs does not affect TH1 responses to systemic or mucosal viral infections.

We have previously reported that global knockout of Bach2 leads to the development of Type 2 cytokine-driven

![Graph](image-url)
fatal eosinophilic crystalline pneumonia [20]. Further, loss of Bach2 in all T cells also leads to aberrant T_{H2} immunity and lung disease [24]. Treg-specific Bach2 deficiency did not lead to the unprovoked development of eosinophilic crystalline pneumonia or T_{H2}-driven lung disease for at least 8-9 months of age. Thus, Bach2 deficiency in Tregs is not sufficient to cause unprovoked T_{H2}-driven lung inflammation. Since variations in the Bach2 gene have been linked to asthma in humans and Bach2 deficiency led to eosinophilic crystalline pneumonia in mice [20, 32, 33], we assessed whether Treg-specific Bach2 deficiency affected the development of fungal protease-induced allergic inflammation in the lungs. Recent work has shown that some allergen proteases possess the ability to cleave and potentiate IL-33 [40, 41]. IL-33 is then capable of driving an ST2-dependent type 2 inflammatory response. Remarkably, despite expressing high basal levels of GATA3 and ST2, Bach2-deficient Tregs appeared to be unable to suppress Aspergillus protease-induced allergic inflammation. Bach2-deficient Tregs expressed a more activated phenotype with protease treatment, and these increases were paralleled by increases of effector phenotypes in WT counterparts as well. Overall, these data suggest that Bach2 is required for Tregs to restrain T_{H2} inflammation. Mechanistically, the absence of Bach2 leads to increased levels of GATA3, BATF, and IRF4, and subsequently more ST2 expression. The inherently higher basal level of an activated/effector tissue Treg phenotype in Bach2-deficient Tregs makes it difficult to detect further activation or display of effector characteristics; however, it should be noted that WT Tregs need to be stimulated in order to reach the basal level of magnitude that Bach2-deficient Tregs express.

5. Conclusions

Maintenance of Treg homeostasis and effector function in lymphoid and nonlymphoid tissues is critical for mitigating autoimmunity and inflammatory diseases. In this manuscript, we ascribe vital roles for Bach2 in regulating the numbers and functions of effector Tregs in lymphoid and non-lymphoid tissues. First, we confirm previous findings [25–27] that Treg-specific Bach2 deficiency leads to unprovoked precocious differentiation of effector Tregs in lymphoid tissues. Second, we document that Bach2 restrains the development of CXCR3+ CD103+ ST2+ tissue Tregs in secondary lymphoid tissues. Third, we show that Bach2 in Tregs limits the number of CXCR3+ CD103+ Tregs in VAT. Fourth, loss of Bach2 in Tregs does not affect Type 1 immunity to systemic and mucosal viral infections. Fifth, we show that Treg-specific Bach2 deficiency does not result in unprovoked T_{H2}-driven inflammation, but restrains aggressive fungus protease-induced Type 2 inflammation in lungs by functioning as a transcriptional checkpoint in both Tregs and conventional effector cells. In summary, findings presented in this manuscript provide mechanistic insights into the role of Bach2 in regulating Treg homeostasis and protecting humans against Type 2 immunity such as asthma and other allergic disorders.

Data Availability

All data presented in this manuscript is archived in secure computers at the University of Wisconsin-Madison.

Conflicts of Interest

Authors have no conflicts of interest with data published in this manuscript.

Authors’ Contributions

AC, DW, JS, BK, and MS. designed, performed, analyzed experiments, and provided conceptual input for the manuscript. WL and BK performed experiments. AC and MS wrote the manuscript, which was proofread by all authors.

Acknowledgments

This work was supported by PHS grants (U01 AI124299 R21 AI149793-01A1) from National Institutes of Health and John E. Butler professorship to M. Suresh. AC was supported by a predoctoral fellowship from the National Institutes of Health (T32AI055397). We gratefully acknowledge Emory NIH Tetramer Core Facility for providing MHC-I and MHC-II tetramers. We would also like to thank genuine appreciation for the efforts of the veterinary and animal care staff at UW-Madison.

References

[1] S. Z. Josefowicz, L. F. Lu, and A. Y. Rudensky, “Regulatory T cells: mechanisms of differentiation and function,” Annual Review of Immunology, vol. 30, no. 1, pp. 531–564, 2012.
[2] K. S. Smigiel, E. Richards, S. Srivastava et al., “CCR7 provides localized access to IL-2 and defines homeostatically distinct regulatory T cell subsets,” The Journal of Experimental Medicine, vol. 211, no. 1, pp. 121–136, 2014.
[3] A. Liston and D. H. Gray, “Homeostatic control of regulatory T cell diversity,” Nature Reviews. Immunology, vol. 14, no. 3, pp. 154–165, 2014.
[4] E. Cretney, A. Xin, W. Shi et al., “The transcription factors Blimp-1 and IRF4 jointly control the differentiation and function of effector regulatory T cells,” Nature Immunology, vol. 12, no. 4, pp. 304–311, 2011.
[5] E. Cretney, A. Kallies, and S. L. Nutt, “Differentiation and function of FOXP3+ effector regulatory T cells,” Trends in Immunology, vol. 34, no. 2, pp. 74–80, 2013.
[6] D. J. Campbell and M. A. Koch, “Phenotypical and functional specialization of FOXP3+ regulatory T cells,” Nature Reviews. Immunology, vol. 11, no. 2, pp. 119–130, 2011.
[7] K. E. Webster, S. Walters, R. E. Kohler et al., “In vivo expansion of T reg cells with IL-2-mAb complexes: induction of resistance to EAE and long-term acceptance of islet allografts without immunosuppression,” The Journal of Experimental Medicine, vol. 206, no. 4, pp. 751–760, 2009.
[8] G. Cheng, A. Yu, M. J. Dee, and T. R. Malek, “IL-2R signaling is essential for functional maturation of regulatory T cells during thymic development,” Journal of Immunology, vol. 190, no. 4, pp. 1567–1575, 2013.
[9] A. Yu, L. Zhu, N. H. Altman, and T. R. Malek, “A low interleukin-2 receptor signaling threshold supports the development and homeostasis of T regulatory cells,” *Immunity*, vol. 30, no. 2, pp. 204–217, 2009.

[10] J. H. Lee, S. G. Kang, and C. H. Kim, “FoxP3+ T cells undergo conventional first switch to lymphoid tissue homing receptors in thymus but accelerated second switch to nonlymphoid tissue homing receptors in secondary lymphoid tissues,” *Journal of Immunology*, vol. 178, no. 1, pp. 301–311, 2007.

[11] M. Feuerer, J. A. Hill, K. Kretschmer, H. von Boehmer, D. Mathis, and C. Benoist, “Genomic definition of multiple ex vivo regulatory T cell subphenotypes,” *Proceedings of the National Academy of Sciences of the United States of America*, vol. 107, no. 13, pp. 5919–5924, 2010.

[12] B. D. Sathre, P. Treuting, N. Perdue et al., “Altering the distribution of Foxp3(+) regulatory T cells results in tissue-specific inflammatory disease,” *The Journal of Experimental Medicine*, vol. 204, no. 6, pp. 1335–1347, 2007.

[13] A. M. Sanchez, J. Zhu, X. Huang, and Y. Yang, “The development and function of memory regulatory T cells after acute viral infections,” *Journal of Immunology*, vol. 189, no. 6, pp. 2805–2814, 2012.

[14] M. Panduro, C. Benoist, and D. Mathis, “Tissue Tregs,” *Annual Review of Immunology*, vol. 34, no. 1, pp. 609–633, 2016.

[15] A. Vasanthakumar, K. Moro, A. Xin et al., “The transcriptional regulators IRF4, BATF and IL-33 orchestrate development and maintenance of adipose tissue-resident regulatory T cells,” *Nature Immunology*, vol. 16, no. 3, pp. 276–285, 2015.

[16] C. Schiering, T. Krausgruber, A. Chomka et al., “The alarmin IL-33 promotes regulatory T-cell function in the intestine,” *Nature*, vol. 513, no. 7519, pp. 564–568, 2014.

[17] B. Afzali, J. Gronholm, V. Vandrovcova et al., “BACH2 immuno-deficiency illustrates an association between super-enhancers and haploinsufficiency,” *Nature Immunology*, vol. 18, no. 7, pp. 813–823, 2017.

[18] R. Roychoudhuri, D. Clever, P. Li et al., “BACH2 regulates CD8+ T cell differentiation by controlling access of AP-1 factors to enhancers,” *Nature Immunology*, vol. 17, no. 7, pp. 851–860, 2016.

[19] S. Tsukumo, M. Unno, A. Muto et al., “Bach2 maintains T cells in a naive state by suppressing effector memory-related genes,” *Proceedings of the National Academy of Sciences of the United States of America*, vol. 110, no. 26, pp. 10735–10740, 2013.

[20] E. H. Kim, D. J. Gasper, S. H. Lee, E. H. Plisch, J. Svaren, and M. Suresh, “Bach2 regulates homeostasis of Foxp3+ regulatory T cells and protects against fatal lung disease in mice,” *Journal of Immunology*, vol. 192, no. 3, pp. 985–995, 2014.

[21] R. Roychoudhuri, K. Hirahara, K. Mousavi et al., “BACH2 represses effector programs to stabilize T(reg)-mediated immune homeostasis,” *Nature*, vol. 498, no. 7455, pp. 506–510, 2013.

[22] L. Yang, S. Chen, Q. Zhao, Y. Sun, and H. Nie, “The critical role of Bach2 in shaping the balance between CD4+ T cell subsets in immune-mediated diseases,” *Mediators of Inflammation*, vol. 2019, Article ID 2609737, 9 pages, 2019.

[23] M. Yamashita and M. Kuwahara, “The critical role of Bach2 in regulating type 2 chronic airway inflammation,” *International Immunology*, vol. 30, no. 9, pp. 397–402, 2018.

[24] M. Kuwahara, W. Ise, M. Ochi et al., “Bach2-Batf interactions control Th2-type immune response by regulating the IL-4 amplification loop,” *Nature Communications*, vol. 7, no. 1, article 12596, 2016.

[25] F. M. Grant, J. Yang, R. Nasrallah et al., “BACH2 drives quiescence and maintenance of resting Treg cells to promote homeostasis and cancer immunosuppression,” *The Journal of Experimental Medicine*, vol. 217, no. 9, 2020.

[26] T. Sidwell, Y. Liao, A. L. Garnham et al., “Attenuation of TCR-induced transcription by Bach2 controls regulatory T cell differentiation and homeostasis,” *Nature Communications*, vol. 11, no. 1, p. 252, 2020.

[27] H. Zhang, D. Dai, Q. Hu et al., “Bach2 attenuates IL-2R signaling to control Treg homeostasis and Tfr development,” *Cell Reports*, vol. 35, no. 6, article 109096, 2021.

[28] Y. P. Rubtsov, J. P. Rasmussen, E. Y. Chi et al., “Regulatory T cell-derived interleukin-10 limits inflammation at environmental interfaces,” *Immunity*, vol. 28, no. 4, pp. 546–558, 2008.

[29] B. M. Matta and H. R. Turnquist, “Expansion of regulatory T cells in vitro and in vivo by IL-33,” *Methods in Molecular Biology*, vol. 1371, pp. 29–41, 2016.

[30] M. Delacher, C. D. Imbusch, A. Hotz-Wagenblatt et al., “Precursors for nonlymphoid-tissue Treg cells reside in secondary lymphoid organs and are programmed by the transcription factor BATF,” *Immunity*, vol. 52, no. 2, pp. 295–312.e11, 2020.

[31] M. Delacher, C. D. Imbusch, D. Weichenhan et al., “Genome-wide DNA-methylation landscape defines specialization of regulatory T cells in tissues,” *Nature Immunology*, vol. 18, no. 10, pp. 1160–1172, 2017.

[32] E. Kreiner, J. Waage, M. Standl et al., “Shared genetic variants suggest common pathways in allergy and autoimmune diseases,” *The Journal of Allergy and Clinical Immunology*, vol. 140, no. 3, pp. 771–781, 2017.

[33] Y. Huo and H. Y. Zhang, “Genetic mechanisms of asthma and the implications for drug repositioning,” *Genes (Basel)*, vol. 9, no. 5, p. 237, 2018.

[34] D. L. Wiesner, R. M. Merkhofer, C. Ober et al., “Club cell TRPV4 serves as a damage sensor driving lung allergic inflammation,” *Cell Host & Microbe*, vol. 27, no. 4, pp. 614–628.e6, 2020.

[35] B. Popovic, M. Golemac, J. Podlech et al., “IL-33/ST2 pathway drives regulatory T cell dependent suppression of liver damage upon cytomegalovirus infection,” *PLoS Pathogens*, vol. 13, no. 4, article e1006345, 2017.

[36] A. Vasanthakumar and A. Kallies, “Interleukin (IL)-33 and the IL-1 family of cytokines-regulators of inflammation and tissue homeostasis,” *Cold Spring Harbor Perspectives in Biology*, vol. 11, no. 3, article a028506, 2019.

[37] L. Guo, G. Wei, J. Zhu et al., “IL-1 family members and STAT activators induce cytokine production by Th2, Th17, and Th1 cells,” *Proceedings of the National Academy of Sciences of the United States of America*, vol. 106, no. 32, pp. 13463–13468, 2009.

[38] Z. Guo, J. Wu, J. Zhao et al., “IL-33 promotes airway remodeling and is a marker of asthma disease severity,” *The Journal of Asthma*, vol. 51, no. 8, pp. 863–869, 2014.

[39] B. M. Matta, J. M. Lott, L. R. Mathews et al., “IL-33 is an unconventional Alarmin that stimulates IL-2 secretion by dendritic cells to selectively expand IL-33R/ST2+ regulatory T cells,” *Journal of Immunology*, vol. 193, no. 8, pp. 4010–4020, 2014.
[40] Y. Hiraishi, S. Yamaguchi, T. Yoshizaki et al., “IL-33, IL-25 and TSLP contribute to development of fungal-associated protease-induced innate-type airway inflammation,” Scientific Reports, vol. 8, no. 1, article 18052, 2018.

[41] I. C. Scott, J. B. Majithiya, C. Sanden et al., "Interleukin-33 is activated by allergen- and necrosis-associated proteolytic activities to regulate its alarmin activity during epithelial damage," Scientific Reports, vol. 8, no. 1, p. 3363, 2018.