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Highlights

Human and mouse ADSCs express IL-33 in response to IL-β stimulation

mADSC-derived IL-33 inhibits inflammation in salivary glands in SS model

mADSC-derived IL-33 expand ST2⁺ Tregs *in vitro* and in SS model

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Adipose-mesenchymal stromal cells suppress experimental Sjögren syndrome by IL-33-driven expansion of ST2⁺ regulatory T cells

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SUMMARY

Adipose-derived mesenchymal stromal cells (ADSCs) play important roles in the alleviation of inflammation and autoimmune diseases. Interleukin-33 (IL-33), a member of the IL-1 family, has been shown to regulate innate and adaptive immunity. However, it is still unknown whether ADSCs regulate immune responses via IL-33. We show here that ADSCs produced IL-33 in response to IL-1 β stimulation, which depended on TAK1, ERK, and p38 pathways. ADSCs-derived IL-33 drove the proliferation of CD4⁺Foxp3⁺ST2⁺ regulatory T cells (Tregs) and alleviated experimental autoimmune Sjögren syndrome in mice. Importantly, human ADSCs also produced IL-33 in response to IL-1 β . Thus, we have revealed a previously unrecognized immunoregulatory function of ADSCs by IL-33 production in experimental autoimmunity, which may have clinical applications for human immunopathology.

INTRODUCTION

Adipose-derived mesenchymal stromal cells (ADSCs), a subtype of mesenchymal stromal cells (MSCs), played important roles in regenerative medicine to differentiating in bone and adipocytes and in modulating immune responses and alleviating inflammation or autoimmune diseases including arthritis, colitis, and autoimmune diabetes (Bassi et al., 2012; Gonzalez et al., 2009a, 2009b; Mizuno et al., 2012; Razmkhah et al., 2015; Shang et al., 2015; Zuk et al., 2001). The MSCs-tracking analysis showed that exogenously delivered MSCs could be detected in inflamed zones for the first week after injection, suggesting their interaction with inflammatory cells and cytokines. Indeed, studies reported that inflammatory cytokines such as interferon (IFN)- γ , interleukin (IL)-1 β , and tumor necrosis factor (TNF)- α could be critical for MSC-mediated immunoregulation (Castelo-Branco et al., 2012; Chen et al., 2013; Tanaka et al., 2008; Xu et al., 2012). The inflammatory cytokines drive the immunosuppression function of MSCs through expression and release of different factors and cytokines in the inflammatory "niches" (Krampera et al., 2006; Meisel et al., 2004; Mougiakakos et al., 2011; Ren et al., 2009; Sheng et al., 2008).

IL-33 was first reported as a novel member of the IL-1 family in 2005 (Schmitz et al., 2005) and is now recognized as a crucial factor in influencing innate and adaptive immunity (Ali et al., 2011). Among its pleiotropic functions (Baumann et al., 2015; Gao et al., 2015; Hepworth et al., 2012; Price et al., 2010; Reichenbach et al., 2015; Yang et al., 2011), IL-33 drives proliferation of CD4⁺Foxp3⁺ regulatory T cells (Tregs) and group 2 innate lymphoid cells (ILC2s) and regulates immune responses in lymphoid organs, gut, lungs, and adipose tissues (Arpaia et al., 2015; Matta et al., 2016; Molofsky et al., 2015b; Mougiakakos et al., 2011; Schiering et al., 2014). IL-33 also functions for the maintenance of tissue homeostasis and repair of tissue damage (Liew et al., 2016; Molofsky et al., 2015a). Typically IL-33 is expressed at high levels in the nuclei of various cell types in human and mouse tissues in the steady state, including endothelial cells (Chen et al., 2015), epithelial cells (Nakanishi et al., 2013), macrophages, and fibroblast-like cells (Moussion et al., 2008; Pichery et al., 2012). Besides, it was recently reported that some subsets of MSCs expressing IL-33 were involved in the homeostasis and metabolism of adipocytes (Mahlakoiv et al., 2019). However, it remains largely unknown what drives MSCs to produce IL-33 and whether the IL-33 has immunoregulatory effects on distant organs and tissues beyond adipose tissues. ¹Mucosal Immunology Section, NIDCR, NIH, Bethesda, MD 20892, USA

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Figure 1. mADSC can produce IL-33 with IL-1 β stimulation

(A) II33 mRNA expression in mADSC stimulated for 6 h with different cytokines as indicated. (2 ng/mL IL-1 β , 10 ng/mL IL-4, 10 ng/mL IL-5, 50 ng/mL IL-6, 10 ng/mL IL-7, 10 ng/mL IL-12, 10 ng/mL IL-13, 10 ng/mL IL-15, 10 ng/mL IL-21, 10 ng/mL IL-22, 10 ng/mL IL-23, 10 ng/mL IL-25, 10 ng/mL IFN γ , 2 ng/mL TGF- β , 10 ng/mL TNF α , 10 ng/mL LPS).

(B) II33 mRNA can be produced in mADSC stimulated for 6 h with 10 ng/mL IL-1 β .

(C) II33 mRNA in mADSC stimulated for 6 h with different dose of concentration of IL-1 β (0, 0.002, 0.02, 0.2, 2 and 20 ng/mL).

(D) Western blot of IL-33 and β -actin in mADSC stimulated with or without 2 ng/mL IL-1 β for 24 or 48 h (E) ELISA of IL-33 in mADSC lysate and supernatant stimulated with or without 10 ng/mL IL-1 β for 72 h. Data are pooled from three (A and C) independent experiments or are representative of three (B, D, and E) independent experiments. In (A), (B), (C), and (E) one-way ANOVA was used. Bars, mean; error bars, SEM; *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.001.

RESULTS

Murine ADSCs produce IL-33 in response to IL-1 β

ADSCs from C57BL/6 mice and IL-33-deficient mice (C57BL/6J-*II33*^{tm1b(EUCOMM)Cln}, *II33^{-/-}*) were isolated from subcutaneous fat tissues. ADSCs were determined based on their positivity for MSC markers SCA-1, CD44, CD90, and CD105 and negativity for epithelial cell marker CD31 and hematopoietic cell marker CD45 (Bourin et al., 2013) (Figure S1A). Also, ADSCs were functionally capable of differentiating into adipocytes, osteoblasts, and chondroblasts under standard differentiation culture conditions (Figure S1B). ADSCs exhibited no difference in their surface markers and multiple differentiation potential from wild-type (WT) and *II33^{-/-}* mice.

To mimic IL-33 production from mADSCs in the inflammatory "niches," we stimulated ADSCs that were isolated from C57BL/6 mice with a panel of inflammatory cytokines in cultures for 6 h *I*/33 mRNA was determined by real-time PCR. Among the comprehensive panel of the cytokines we examined, only IL-1 β was able to significantly upregulate *I*/33 mRNA expression in ADSCs (Figure 1A). ADSCs expressed





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Figure 2. Continued

(D) Western blot of IL-33 and GAPDH in mADSC stimulated with or without IL-1 β , TAK1 inhibitor, ERK inhibitor, or p38 inhibitor for 24 h. (E) Western blot of p-ERK, ERK, p-p38, p38, and GAPDH in mADSC stimulated with or without IL-1 β , TAK1 inhibitor for 1 h. Data are pooled from three (B and C) independent experiments or are representative of three (A, D, and E) independent experiments. In (A), Student's t test was used. In (B), (C), and (E) one-way ANOVA was used. Bars, mean; error bars, SEM; *p < 0.05, **p < 0.01.

spontaneous base levels of *II33* mRNA higher than did T cells (Figure 1B). IL-1 β driving *II33* mRNA expression in ADSCs was dose-dependent and reached a plateau at 0.2 ng/mL of IL-1 β (Figure 1C). Consistent with *II33* mRNA, western blot analysis showed a substantially higher level of IL-33 protein in IL-1 β -treated ADSCs compared with untreated cells (Figure 1D). ELISA analysis also showed higher levels of IL-33 protein in the cell lysates from IL-1 β -treated ADSCs when compared with the untreated cells, although IL-33 was hardly detected in the cell culture supernatants from all the cultures (Figure 1E). Of note, although IFN- γ alone was unable to induce *II33* mRNA, it enhanced IL- β -induced IL-33 in mouse ADSCs (Figure S2). Thus, mADSCs produce IL-33 upon IL-1 β stimulation.

IL-1β upregulates IL-33 through TAK1-ERK/p38 pathways in mADSCs

We next investigated the signaling pathways by which IL-33 was induced by IL-1 β stimulation in ADSCs. We first analyzed the expression and activation of ERK, p38, JNK, and nuclear factor (NF)- κ B proteins in ADSCs stimulated with IL-1 β . We found that the relative ratio of phosphorylated ERK (p-ERK)/ERK and phosphorylated p38 (p-p38)/p38 were increased in ADSCs treated with IL-1 β compared with the untreated cells (Figure 2A). However, there were no significant differences between the phosphorylated JNK and NF- κ B (p65) proteins between IL-1 β -treated and untreated ADSCs (Figures S3A and S3B).

Next, we investigated which of the aforementioned pathways was involved in IL-1 β -driven IL-33 induction in ADSCs by utilizing the selective protein kinase inhibitors 0.2 μ M U0126 (for ERK), 20 μ M SB203580 (for p38 MAPK), 2 μ M SP600125 (for JNK), and 2 μ M JSH23 (for NF- κ B) to block the activity of the indicated proteins. We found that inhibition of ERK and p38 MAPK activity significantly reduced IL-1 β -induced *II33* mRNA and IL-33 protein (Figures 2B and 2D), whereas inhibition of JNK and NF- κ B failed to do so (Figures S3C and S3D). In addition, suppression of TAK1 activity also decreased *II33* mRNA and IL-33 protein in ADSCs (Figures 2C and 2D). To explore the mechanism of the TAK1 signal pathway during IL-1 β -driven IL-33, we examined whether TAK1 inhibition affects the phosphorylation of p38 and ERK. Our result showed that the phosphorylation of p38 and ERK was significantly reduced in TAK1 inhibitor (100 nM) group after IL-1 β treatment (Figure 2E). These data suggest that IL-1 β activates TAK1, up-regulating phosphorylation of p38 and ERK, and subsequently mediates IL-33 expression.

mADSCs-derived IL-33 regulates ST2 expression in Tregs in vitro

To determine the immunoregulatory function of IL-33 produced by ADSCs, we next studied the effects of IL-1 β -treated mADSCs lysates on CD4⁺CD25⁺Foxp3⁺ Tregs (Chen et al., 2003), because IL-33 has been shown to promote the proliferation of Tregs (Matta et al., 2016; Vasanthakumar et al., 2015). ADSCs from WT or *II33^{-/-}* mice were cultured with or without IL-1 β to confirm that *II33* mRNA and IL-33 protein were detected only in WT but not *II33^{-/-}* cells (Figures 3A and 3B). CD4⁺CD25⁺ Tregs isolated from spleen and lymph nodes in C57BL/6 mice were cultured with IL-1 β -stimulated WT or *II33^{-/-}* ADSCs lysates. We found that IL-1 β -stimulated WT ADSCs lysates significantly upregulated *II1r*/1 mRNA (receptor for IL-33, ST2) expression in Tregs, which was abrogated by the inclusion of anti-IL-33 antibody in the cultures (Figure 3C). Consistently, IL-1 β -stimulated *II33^{-/-}* ADSCs lysates exhibited significantly lower activity to upregulate *II1r*/1 mRNA in Tregs than did WT ADSCs lysates (Figure 3C). Consequently, the frequency and the absolute number of ST2⁺ Treg subsets within the CD4⁺Foxp3⁺ Tregs cultured with IL-1 β -treated WT ADSCs were much higher than those of ST2⁺ Tregs in Tregs stimulated with *WT* ADSCs was entirely abolished by neutralization of IL-33 with anti-IL33 antibody (Figure 3E). The data collectively indicate that IL-33 from mADSCs upregulates ST2⁺ Tregs in cultures.

WT but not *II33^{-/-}* mADSCs improve Sjögren syndrome by suppressing inflammation in salivary glands in NOD mice

The expansion of ST2⁺ Tregs by ADSCs-derived IL-33 encouraged us to investigate the immunoregulatory and therapeutic function of ADSCs in autoimmunity and inflammation *in vivo*. For this, we intravenously injected WT and $I/33^{-/-}$ mADSCs (both on C57BL/6 background) into allogeneic NOD/ShiLtj mice (NOD, on







Figure 3. IL-33 from mADSC affects ST2 expression on Treg in vitro

(A) II33 mRNA expression in ADSC from WT and II33^{-/-} mice stimulated for 6 h with or without 10 ng/mL IL-1 β . (B) ELISA of IL-33 protein in mADSC stimulated with 10 ng/mL IL-1 β for 72 h

(C) II1rl1 mRNA expression in CD4⁺CD25⁺ T cell from spleen with anti-CD3/CD28 plus stimulated with or without ADSC lysate from WT or II33^{-/-} mice or ADSC soap from WT mice with anti-IL-33 for 24 h.

(D) Frequency and absolute number of Foxp 3^+ T cells in CD4⁺CD25⁺T cell from spleen with anti-CD3/CD28 plus stimulated with or without ADSC lysate from WT or $II33^{-/-}$ mice or ADSC soap from WT mice with anti-IL-33 for 72 h.





Figure 3. Continued

(E) Frequency and absolute number of ST2⁺ T cells in CD4⁺Foxp3⁺ T cell from spleen with anti-CD3/CD28 plus stimulated with or without ADSC lysate from WT or *II33^{-/-}* mice or ADSC soap from WT mice with anti-IL33 for 72 h. Data are pooled from three (A, C, D, and E) independent experiments or are representative of three (B) independent experiments. In (B), Student's t test was used. In (A), (C), (D), and (E), one-way ANOVA was used. Bars, mean; error bars, SEM; *p < 0.05, **p < 0.01, ***p < 0.001, ***p < 0.001.

Cataract Shionogi background) that have been well known to develop an autoimmune Sjögren syndrome (SS) spontaneously. The rationale underlying the utilization of allogeneic ADSC transplantation was based on the published animal and clinical studies that show negligible allorejection and side effects of allogeneic MSCs transplantation due to the non/low MHC II expression on MSCs, yet exhibition of beneficial and immunoregulatory function (Alexeev et al., 2014; Xu et al., 2012). We injected allogeneic ADSCs from WT or I/33^{-/-} mice or PBS into NOD/ShiLtj mice at 7 weeks of age when the inflammation starts to occur in the submandibular glands. Yet, the secretory function of the submandibular glands has not been compromised (Delaleu et al., 2011). The saliva flow rates were measured weekly from 7 through 13 weeks (Figure S4A). We found that WT ADSCs transplantation significantly improved the saliva flow rates 2 weeks after injection in NOD/ShiLtj mice compared with PBS- or I/33-/- mADSCs-treated mice (Figure 4A). We then examined the expression of genes, including Slc12a2, Tjp1, Agp5, Itpr3, Trpv4, and Chrm3, which have been known to be relevant to submandibular gland function at the 13th week (Markadieu and Delpire, 2014; Wang et al., 2017). The expressions of all those genes except Agp5 mRNA in submandibular glands were significantly increased in WT ADSCs-treated mice compared with the mice treated with I/33^{-/-} mADSCs or PBS (Figures 4B-4G). Consistent with the improvement of saliva flow and the upregulation of the genes associated with submandibular gland function, histological analysis of the submandibular gland tissues revealed that transplantation of WT ADSCs resulted in a substantial reduction of inflammatory areas in the submandibular glands compared with $II33^{-/-}$ ADSCs- or PBS-treated groups (Figures 4H and 4I). These findings altogether indicate that allogeneic ADSCs suppress inflammation and improve the secretory function of salivary glands in an IL-33-dependent manner.

ADSC-derived IL-33 upregulates ST2⁺ Tregs in NOD mice

To understand the underlying mechanisms by which allogeneic ADSCs IL-33 suppressed inflammation and improved the function of submandibular glands, we examined the changes of Tregs and proinflammatory effector cells in the submandibular glands of NOD mice. Strikingly, WT ADSCs treatment resulted in a significant increase in both the frequency and absolute number of $CD4^+Foxp3^+$ Tregs in the submandibular glands in SS-like NOD mice compared with other groups (Figure 5A). More importantly, WT ADSCs treatment, but not $II33^{-/-}$ ADSCs treatment, led to a significant increase in the frequency and total number of ST2⁺ Tregs (Figure 5B). Further analysis of *in vivo* Treg expansion with Ki67 staining revealed that WT ADSCs treatment caused significantly more Ki67⁺ Tregs in the submandibular glands than did $II33^{-/-}$ ADSCs or PBS treatment, suggesting IL-33-driven Treg proliferation upon WT ADSCs transplantation (Figure 5C).

As IL-33 is also associated with the differentiation and function of various lymphocytes including type 2 helper T (Th2) cells and ILC2s (Licona-Limon et al., 2013; Molofsky et al., 2013; Moro et al., 2010; Schmitz et al., 2005) in addition to Tregs, we studied Th2 (IL-4⁺, IL-13⁺) CD4⁺ T cells and ILC2s in the submandibular glands among all the groups of NOD mice and no significant differences could be observed (Figures S4–S7). Furthermore, the frequency of IL-13-expressing (Lin⁻CD45⁺GATA3⁺ST2⁺IL-13⁺) and IL-4-producing (Lin⁻CD45⁺GATA3⁺ST2⁺IL-4⁺) ILC2 subsets also did not exhibit changes after ADSCs treatment (Figures S8A–S8C). Interestingly, the frequency of CD8⁺TCR $\alpha\beta^+$ IL-4⁺ T cells increased in the submandibular glands of WT ADSCs-treated group compared with *II33^{-/-}* ADSCs- or PBS-treated groups (Figure S6D), but the significance of the increase remains unknown.

We next examined the CD45⁺ inflammatory cells in the submandibular glands. WT ADSCs treatment caused significantly fewer CD45⁺ cells in the submandibular glands than *II33^{-/-}* ADSCs or PBS-treated mice (Figure 5D). The absolute number of CD4⁺ (Figure 5E) and CD8⁺ (Figure 5F) T cells, but not TCR $\gamma\delta^+$ T cells (Figure 5G), significantly decreased in WT ADSC-treated mice compared with *III33^{-/-}* ADSCs- or PBS- treated groups. Consequently, the total number of IFN- γ^+ CD4⁺ T cells (Th1) and IFN- γ^+ CD8⁺ T cells were decreased in WT ADSC-treated NOD mice, although the frequency did not change significantly (Figures S4B–S4E, S5A, S5E, S6A, S6E, S7A, and S7E). Interestingly, both WT and *II33^{-/-}* mADSCs treatment showed a similar reduction of IL-17⁺ T cells compared with the PBS group, suggesting that









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Figure 4. WT but not *II33^{-/-}* **mADSCs suppress inflamed tissue damage and improve submandibular gland function in NOD/ShiLtj mice** (A) Salivary flow rate of NOD/ShiLtj treated with PBS, ADSC from WT or *II33^{-/-}* mice. PBS group (n = 12), KO group (n = 12), WT group (n = 12). (B) Slc12a2 mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. (C) *Tjp1* mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. (D) *Itpr3* mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.





Figure 4. Continued

(E) *Trpv4* mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.
(F) *Chrm3* mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.
(G) *Aqp5* mRNA expression in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.
(H) Histology of the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.
(I) The infiltrating area in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.
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the decrease of IL-17 was independent of IL-33 (Figures S5B and S7B). In contrast to the submandibular glands, there was no significant difference in the frequency and number of CD4⁺Foxp3⁺ Tregs, ST2⁺Tregs and Ki67⁺ Tregs in the spleen, and armpit, inguinal and submandibular lymph nodes of all groups of NOD/ ShiLtj mice (Figure S9). Taken together, these data indicate that mADSCs-IL-33-mediated suppression of inflammation and improvement of submandibular gland function was mainly ascribed to the enhanced number of ST2⁺ Tregs and reduced the number of CD45⁺ proinflammatory cells especially IFN- γ^+ T cells in the glands.

IL-1β upregulates IL-33 in human ADSCs

We next investigated whether human ADSCs also expressed IL-33 and IL-1 β upregulated its expression. For this, we cultured human ADSCs with or without 2 ng/mL IL-1 β for 6 and 24 h and then checked *IL*33 mRNA and IL-33 protein, respectively. Untreated human ADSCs hardly exhibited IL-33; however, IL-1 β treatment significantly upregulated the amount of *IL*33 mRNA in human ADSCs at 6 h (Figure 6A). IL-33 protein was also increased significantly by IL-1 β treatment in human ADSCs at 24 h (Figure 6A). IL-33 protein was also increased significantly by IL-1 β treatment in human ADSCs at 24 h (Figure 6B). We also studied the signal pathways for IL-33 production by human ADSCs in response to IL-1 β . Consistent with murine ADSCs, the levels of phosphorylated ERK and p38 proteins were significantly increased in human ADSCs after IL-1 β stimulation compared with the unstimulated cells (Figure 6C). Moreover, the selective protein kinase inhibitors (5Z)-7-oxozeaenol, U0126, and SB203580, which inhibit TAK1, ERK, and p38 MAPK, respectively, significantly decreased the inductive effect of IL-1 β on *IL33* mRNA and IL-33 protein in human ADSCs (Figures 6D and 6E). TAK1 inhibitor also decreased the relative ratio of phosphorylated ERK/ERK and phosphorylated p38/p38 (Figure 6F). These data indicated that human ADSCs also produce IL-33 by IL-1 β stimulation through TAK1-ERK/p38 signal pathways, suggesting a translational relevance and significance of the murine ADSCs treatment in animal SS models.

DISCUSSION

In this article, we demonstrated that ADSCs represent an important cellular source of IL-33 that could be induced specifically by IL-1 β among the dozens of immune cytokines tested. We further elucidated that IL-1 β -induced IL-33 production by ADSCs was mediated by TAK1-ERK/p38 signaling pathways. By using $I/33^{-/-}$ mice, we have uncovered that allogeneic ADSCs transplantation was able to suppress the inflammation and improve the saliva secretory function of salivary glands in NOD mice via an IL-33-dependent manner. Importantly, we have elucidated that the immunoregulatory function in SS by ADSCs-derived IL-33 is mediated mainly through upregulation of ST2⁺ Tregs in the submandibular glands. Strikingly, we have shown that human ADSCs also produce IL-33 in response to IL-1 β stimulation *in vitro*, suggesting a clinical relevance and significance in human diseases such as SS.

Several conclusions can be drawn from our findings. First, ADSCs are an important cellular source of IL-33 and IL-1 β is the most important cytokine to induce IL-33 in ADSCs. IL-33 has been recognized as a unique but crucial regulatory cytokine in immunoregulation (Baumann et al., 2015; Gao et al., 2015; Hepworth et al., 2012; Price et al., 2010; Reichenbach et al., 2015; Yang et al., 2011) in several physiological and pathological settings such as autoimmunity, obesity, and allergy/asthma (Nakanishi et al., 2013; Nechama et al., 2018; Scott et al., 2018; Zhu et al., 2017). Several types of cells have been reported to produce IL-33 including endothelial cells (Chen et al., 2015), epithelial cells (Nakanishi et al., 2012) (Su et al., 2013; Xu et al., 2008) in response to certain stimuli such as LPS, IFN- γ , TNF- α , IL-3, IL-4, IL-17, and IL-1 β (Meephansan et al., 2012, 2013; Su et al., 2013; Xu et al., 2008; Zhao and Hu, 2012). However, it was unknown whether MSCs could produce IL-33 until recently. Few studies have started to show that MSCs can produce IL-33 (Mahlakoiv et al., 2019), but what is the factor(s)/cytokines to stimulate MSCs to produce IL-33 remained unknown. We have here extended the studies by unambiguously proving that ADSCs indeed produce IL-33. Importantly, we have discovered that IL-1 β is the critical factor in driving IL-33 expression in



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Figure 5. WT ADSCs increase the frequency and an absolute number of CD4⁺Foxp3⁺ST2⁺ Tregs in the submandibular glands of NOD/ShiLtj mice (A) Frequency and absolute number of Foxp3⁺ T cells in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. (B) Frequency and absolute number of ST2⁺ T cells in CD4⁺Foxp3⁺ T cell in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(C) Frequency and absolute number of Ki67⁺ in CD4⁺Foxp3⁺ T cells in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or II33^{-/-} mice.

(D) Absolute number of CD45⁺ cells in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(E) Absolute number of CD45⁺TCR β ⁺CD4⁺ cells per mg of submandibular gland of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. (F) Absolute number of CD45⁺TCR β ⁺CD8⁺ cells in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. (G) Absolute number of CD45⁺TCR γ ^{8⁺} cells in the submandibular glands of NOD/ShiLtj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. Data are

pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; *p < 0.05, **p < 0.01, ***p < 0.001, ***p < 0.001.

ADSCs. This finding is significant as it provides a previously unrecognized mechanism underlying MSCsmediated immunoregulatory activities in the inflammatory tissues and organs and the paradoxical requirement of proinflammatory cytokines such as IL-1 β (Bassi et al., 2012; Ding et al., 2010; Liu et al., 2013; Meisel et al., 2004). The specific stimulatory function for IL-1 β in MSC-IL-33 production is unexpected but exciting. Although it is presently unknown why IL-1 β , but not other proinflammatory cytokines such as TNF- α and IL- δ , is the key to induce IL-33 secretion by ADSCs, it does suggest a possible functional link between IL-1 β -producing cells and ADSCs. As IL-1 β can be a cell membrane-bound form, this finding also suggests an interaction and functional regulation via a cell-cell contact manner, a fascinating question to be explored. The specific role of IL-1 β in inducing IL-33 by ADSCs is further supported by the downstream molecular pathways that are involved in the effect, which is mainly mediated by ERK and p38 and TAK1, rather than NF- κ B and JNK. As shown by qPCR analysis in Figures 1 and 5, we observed that human ADSCs expressed higher levels of IL-33 than did mouse ADSCs in response to their respective IL-1 β stimulation. Although mechanistically still elusive, the species difference was reported before. For example, a study reported that human IL-33 was converted into more stable mature forms; however, mouse IL-33 was rapidly degraded (Cayrol et al., 2018). Thus further work is needed to understand the specific variation in the future.

Second, ADSCs can regulate inflammation and suppress autoimmunity in distant tissues and organs by producing IL-33. Supporting this conclusion includes that administration by intravenous injection of ADSCs from WT mice, but not from $I/33^{-/-}$ mice, effectively suppressed the ongoing inflammation in the submandibular glands and protected the secretary function of saliva in NOD mice. Our findings here provided a mechanism for the immunoregulatory effect of allogenic ADSCs through an IL-33-dependent mechanism.

Moreover, we found that the immunoregulatory function of ADSC-derived IL-33 is mainly through the expansion of ST2⁺ Treqs in the submandibular glands. Several pieces of evidence support this conclusion: *in vitro*, WT, but not I/33^{-/-}, ADSCs-mediated expansion of Tregs are primarily restricted to the ST2⁺ Tregs. In contrast, ADSCs fail to convert naive CD4⁺ T cells to Foxp3⁺ Tregs, consistent with the notion that TGF-B (Chen et al., 2003) rather than IL-33 is the primary driving force to induce Foxp3 from naive CD4⁺ T cells. IL-33 is the cytokine to expand ST2⁺ Tregs. This IL-33-driven ST2⁺ Treg expansion was confirmed in the submandibular glands in SS NOD mice. Administration of WT, but not $I/33^{-/-}$, ADSCs significantly expand ST2⁺ Tregs in the submandibular glands of SS mice, which consequently protects the function of salivary glands. Notably, the increase in ST2⁺ Tregs in the submandibular glands is not only restricted to the frequency but also the absolute number of the ST2⁺ Tregs, further confirming the real growth of the Tregs by IL-33. The significant expansion of ST2⁺ Tregs in the submandibular glands plays an important role in suppressing the inflammation of the tissue. This is supported by the significant decrease in the absolute number of infiltrated CD45⁺ proinflammatory cells in the submandibular glands of SS-like NOD mice treated with WT ADSCs. IL-13-producing T cells have been implicated in the pathogenesis of an SS animal model that is induced by the deficiency of transcriptional factor Id3 (Li et al., 2004). However, we did not find that IL-13⁺ T cells were changed by ADSCs treatment, suggesting the ADSCs-IL-33 functioning NOD model was not primarily by affecting IL-13⁺ Th2 cells. Of note, WT ADSCs treatment surprisingly increased the frequency and number of CD8⁺IL-4⁺ T cells in the submandibular glands; although the overall frequency is low, it remains to be known whether these $IL-4^+$ T cells play any function in the immunoregulatory effects on SS. The suppressive function of Tregs and ST2⁺ Tregs in immune cells and responses has been validated in many in vitro and in vivo settings in the literature, including in NOD mice (Alunno et al., 2015; Burzyn et al., 2013; Katsifis et al., 2007; Sarigul et al., 2010).

Third, IL-33 can be produced by ADSC with IL-1 β stimulation, although IL-33 but cannot be secreted. Supporting this conclusion includes that IL-33 was hardly detected in the cell culture supernatants, although

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Figure 6. IL-1β upregulates IL-33 expression in human ADSCs

(A) IL33 mRNA expression in human ADSC stimulated for 6 h with or without 2 ng/mL IL-1β.

(B) Western blot of IL-33 and GAPDH in human ADSC stimulated with or without 2 ng/mL IL-1 β for 24 h.

(C) Western blot of p-ERK, ERK, p-p38, p38, and GAPDH in human ADSC stimulated with or without IL-1β for 2 h

(D) *IL33* mRNA expression in human ADSC stimulated for 6 h with or without IL-1 β , TAK1 inhibitor, ERK inhibitor, or p38 inhibitor for 6 h.

(E) Western blot of IL-33 and GAPDH in human ADSC stimulated with or without IL-1 β , TAK1 inhibitor, ERK inhibitor, or p38 inhibitor for 24 h.

(F) Western blot of p-ERK, ERK, p-p38, p38, and GAPDH in human ADSC stimulated with or without IL-1 β , TAK1 inhibitor for 1 h. Data are pooled from three (A and D) independent experiments or are representative of three (B, C, E and F) independent experiments. In (A), Student's t test was used. In (D), one-way ANOVA was used. Bars, mean; error bars, SEM; **p < 0.01, ****p < 0.001.





ELISA analysis also showed higher levels of IL-33 protein in the cell lysates from IL-1 β -treated ADSCs compared with the untreated cells. Moreover, the cell lysates from IL-1 β -stimulated WT ADSCs significantly upregulated *ll1rl1* mRNA (receptor for IL-33, ST2) expression in Tregs. There must be a way for MSCs *in vivo* to be equipped functionally and recruited accurately to damaged tissue. Several researches have proved that following intravenous infusion of MSC populations expanded *in vitro*, almost 80% MSCs are trapped in the lungs, and these cells in lung disappeared with a half-life of about 24 h; some MSCs subsequently become home to damaged tissue (Barbash et al., 2003; Lee et al., 2009). We have also previously reported that allogenic MSCs could exhibit protective effects on SS development in mice by showing that MSCs could migrate into the submandibular glands to function (Xu et al., 2012). Therefore, the therapeutic effects of MSCs may depend mainly on the release of IL-33 of dead MSCs to regulate inflammation and tissue homeostasis.

Finally, the ADSCs-IL-33 may also play an important role in MSC-mediated immunoregulation in relevant human autoimmune diseases like SS. Supporting this conclusion include our findings that human ADSCs also produce IL-33, which is also triggered by IL-1 β stimulation *in vitro*. Although it still needs to be confirmed in the patients, our experimental evidence indeed shows the clinical relevance of our animal studies.

In sum, we have discovered that adipose tissues contain MSCs that produce IL-33 in response to proinflammatory cytokine IL-1 β . This MSC-derived IL-33 can suppress inflammation in organ-specific autoimmunity such as SS through specifically expanding ST2⁺ Tregs. Our findings may be clinically and translationally relevant, as human ADSCs also produce IL-33 in response to IL-1 β . These findings collectively should have implications for better manipulation of allogeneic MSCs in relevant human autoimmune diseases such as SS.

Limitations of the study

There are still some limitations in our study. For example, it remains to be elucidated how important IL- 1β -mediated increase in IL-33 from ADSC is in the SS model. The technical challenge and difficulty prevented us to use ST2 knockout Tregs to provide further evidence of IL-33/ST2 pathway in Treg-mediated suppression of SS models, which we will address in the future. Moreover, it would be interesting to investigate whether ADSC-derived IL-33 has any effects on Treg function through their ST2 receptors in patients with SS.

Resource availability

Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Wanjun Chen(wchen@dir.nidcr.nih.gov).

Materials availability

This study did not generate new unique reagents.

Data and code accessibility

This study did not generate/analyze datasets and code.

METHODS

All methods can be found in the accompanying Transparent methods supplemental file.

SUPPLEMENTAL INFORMATION

Supplemental information can be found online at https://doi.org/10.1016/j.isci.2021.102446.

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AUTHOR CONTRIBUTIONS

O.L. and J.X. designed and performed experiments, analyzed and interpreted the data, and drafted the manuscript; F.W., W.J., N.G., P.Z., N.G., Y.H., and A.B. performed experiments; G.M., S.W., and Z. T. provided critical input and/or support. W.C. conceived and supervised the whole study, designed the experiments, and wrote the manuscript.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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Supplemental information

Adipose-mesenchymal stromal cells suppress

experimental Sjögren syndrome by IL-33-driven

expansion of ST2⁺ regulatory T cells

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Figure S1. Characterization of the immunophenotype and differentiation potential of mADSC from C57BL/6 or C57BL/6J-*II33^{tm1b(EUCOMM)Cln*</sub> in vitro. Related to Figure 1. (A) mADSCs from C57BL/6 or C57BL/6J-*II33^{tm1b(EUCOMM)Cln*} express MSC markers. (B) mADSCs from C57BL/6 or C57BL/6J-*II33^{tm1b(EUCOMM)Cln*} were capable of differentiating into adipocytes,}

osteoblasts, and chondroblast under inductive culture conditions.

Data are pooled from three (A,B) independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; *****P* < 0.0001.



Figure S2. Expression pattern of IL-33 induced by IL-1 β synergistically work with TNF- α and IFN- γ . Related Figure 2.

II33 mRNA expression in mADSC stimulated with indicated combinations of recombinant cytokines (10 ng/ml each) for 6 hr.

Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; *P< 0.05, **P < 0.01.



Figure S3. IL-1 β upregulates IL-33 expression in mADSC independent on JNK and NF- κ B(p65) pathways. Related to Figure 2.

(A) Western blot of p-JNK, JNK, and GAPDH in m ADSC stimulated with or without 2ng/ml IL-1 β for 2 hr. (B) Western blot of p-NF-kB(p65), NF-kB(p65) and GAPDH in m ADSC stimulated with or without IL-1 β for 2 hr.

(C) *II33* mRNA expression in mADSC stimulated for 6 hr with or without 2ng/ml IL-1 β plus indicated the concentration of SP600125.

(D) *II33* mRNA expression in mADSC stimulated for 6 hr with or without 2ng/ml IL-1 β plus indicated the concentration of JSH23.

Data are pooled from three (C and D) independent experiments or are representative of three (A and B) independent experiments. In (C and D), one-way ANOVA was used. Bars, mean; error bars, SEM.



Figure S4. CD45⁺ immune cells express in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. Related to Figure 4.

(A) Experimental workflow of allogeneic mADSCs from WT or il33^{-/-} mice, or PBS infused to NOD/Ltj mice at 7 weeks of age.

(B) Frequency of CD45⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(C) Frequency of CD45⁺TCR β ⁺CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(D) Frequency of CD45⁺ TCR β ⁺CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(E) Frequency of CD45⁺ TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM.



Figure S5. Cytokine expression in TCR β^+ CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with ADSCs from WT or *II33^{-/-}* mice. Related to Figure 5.

(Å) Frequency and absolute numbers of IFN γ^+ in TCR β^+ CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(B) Frequency and absolute numbers of IL-17⁺ in TCR β ⁺CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33*^{-/-} mice.

(C) Frequency and absolute numbers of IL-13⁺ in TCR β ⁺CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33*^{-/-} mice.

(D) Frequency and absolute numbers of IL-4⁺ in TCR β ⁺CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33*^{-/-} mice.

(E) Frequency and absolute numbers of TNF- α^+ in TCR β^+ CD4⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; **P*< 0.05.



Figure S6. Cytokine expression in TCR β^+ CD8⁺ cells in the submandibular glands of NOD/Ltj mice treated with ADSCs from WT or *II33^{-/-}* mice. Related to Figure 5.

(A) Frequency of IFN γ^+ in TCR β^+ CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(B) Frequency of IL17⁺ in TCR β^+ CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(C) Frequency of IL13⁺ in TCR β ⁺CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(D) Frequency of IL4⁺ in TCR β ⁺CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(E) Frequency TNF α^+ in TCR β^+ CD8⁺ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; **P*< 0.05.



Figure S7. Cytokine expression in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice treated with ADSCs from WT or *II33^{-/-}* mice. Related to Figure 5.

(A) Frequency of IFN γ^+ in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(B) Frequency of IL17⁺ in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33*^{-/-} mice.

(C) Frequency of IL13⁺ in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

(D) Frequency of IL4⁺ in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(E) Frequency of TNF α^+ in TCR $\gamma\delta^+$ cells in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice. Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM; ***P*< 0.01.



Figure S8. ST2, IL-13, and IL-4 express in ILC2 in the submandibular glands of NOD/Ltj mice injected with ADSCs from WT or *II33^{-/-}* mice. Related to Figure 5.

(A)The frequency of ST2⁺ in ILC2 in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(B) The frequency of IL-13⁺ in ILC2 in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(C) The frequency of IL-4⁺ in ILC2 in the submandibular glands of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

Data are pooled from three independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM.





(A) Frequency of CD4⁺Foxp3⁺ T, ST2 in Treg and Ki67 in Treg cells in the submandibular lymph node of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(B) Frequency of CD4⁺Foxp3⁺ T, ST2 in Treg and Ki67 in Treg cells in the spleen of NOD/Ltj mice injected with PBS, ADSC from WT or *II33^{-/-}* mice.

(C) Frequency of CD4⁺Foxp3⁺ T, ST2 in Treg and Ki67 in Treg cells in the armpit and inguinal lymph nodes of NOD/Ltj mice injected with PBS, ADSC from WT or *II*33^{-/-} mice.

Data (A) are pooled from three independent experiments. Data (B and C) are pooled from two independent experiments. One-way ANOVA was used. Bars, mean; error bars, SEM.

Transparent Methods

Mice

Male mice *C57BL/6*, Female NOD/ShiLtj mice (*Cdh23ahl*) mice were purchased from The Jackson Laboratory. Male *mice C57BL/6J- II33^{tm1b(EUCOMM)Cln*} were donated by Washington University at St. Louis. All animal studies were approved by the Animal Care and Use Committees of National Institute of Dental and Craniofacial Research, National Institutes of Health. 6 to10 weeks old mice were used for all experiments.

Mice/human ADSCs culture

ADSCs were prepared from *C57BL/6* or *C57BL/6J-II33*^{tm1b(EUCOMM)CIn} subcutaneous fat as described previously⁽¹⁾ and cultured for 2 weeks. Human ADSCs were purchased from Lonza Walkersville Inc, which were isolated from healthy (non-diabetic) adult lipoaspirates collected during elective surgical liposuction procedures. Mice and human ADSCs were cultured in alpha-minimum essential medium containing 20% fetal calf serum and 1% antibiotic-antimycotic (Gibco) in a 5%CO₂ incubator at 37°C. In brief, tissue was minced with two scalpels (crossed blades) and then incubated in a 0.5% collagenase l/phosphate buffered saline (PBS) solution (Collagenase Type: CLS, Biochrom, Berlin, Germany; PBS, Sigma, Taufkirchen, Germany) for 40min at 37°C under constant shaking. The digested tissue solution was then separated through a 100 µm strainer, and the resulting filtrate was centrifuged at 300 g for 5 min. The resulting pellet was washed twice with medium and centrifuged again at 300 g for 5 min. Finally, cells were plated for initial cell culture and cultured at 37°C in an atmosphere of 5% CO₂ in the humid air. alpha-minimum essential medium (α -MEM, Gibco) with a physiologic glucose concentration (100 mg/dL) and used as a standard culture medium. The medium was replaced every three days. Subconfluent cells (90%) were passaged by trypsinization. Cells between passages 2 and 3 were used throughout the experiments. Cell morphology was examined by phase contrast microscopy.

CD4⁺CD25⁺Treg isolation and culture

CD4⁺CD25⁺Tregs from spleen and lymph node from C57BL/6 mice were purified by magnetic cell sorting (Miltenyi Biotec). Tregs were cultured in completed DMEM with or without ADSCs lysates.

ADSCs lysates preparation

ADSCs from WT or *II33^{-/-}* mice were cultured with IL-1β for 72h. Then ADSCs were broken with ultrasonic method (output power 500W, twice, 5min/each, 4°C) and centrifuged at 14000rpm for 20min.Then the supernatant was collected. CD4⁺CD25⁺Tregs were cultured with IL-1β-stimulated WT or *II33^{-/-}* ADSC lysates for 24h and 72h with indicated concentration of plate-bound 1µg/ml anti-CD3, 1µg/ml soluble anti-CD28 and 100U/ml IL-2 for 3 days. *II1rl1* mRNA in Tregs and ST2⁺Treg were detected by real-time PCR and flow cytometry analysis.

Real-time PCR

Total RNA was derived from cells using RNeasy Mini Kit (QIAGEN) and reversed transcribed using High Capacity cDNA Reverse Transcription Kit (Applied Biosystems). Quantitative real-time PCR was performed according to the protocol of TaqMan Gene Expression Master Mix (Applied Biosystems) with the following TaqMan primers: Hprt, Mm00446968_m1; MmII33, Mm00505403_m1; MmSt2, Mm00516117_m1; MmCebpα, Mm00514283_s1; MmFabp4, Mm00445878_m1; MmPpar, Mm01184322_m1; MmAlp, Mm00475834_m1; MmSox9, Mm00448840_m1; MmGapdh, Mm99999915_g1; MmFoxp3, Mm00475162_m1; Human Gapdh Hs02758991_m1; Human II33, Hs04931857_m1.

Western blot analysis

Protein was extracted using RIPA buffer containing proteinase inhibitor cocktail and quantified with a Bio-Rad protein assay. An equal amount of protein was separated on SDS-polyacrylamide gels in a Tris/SDS buffer system and then transferred onto nitrocellulose membranes. Blotting was performed according to standard procedures with primary antibodies against IL-33, ERK, p38, JNK, NF- κ B(p65), p-ERK, p-p38, p-JNK, p-NF- κ B (p65), β -actin and GAPDH overnight, followed by incubation with appropriate fluorescenceconjugated secondary antibodies. The proteins of interest were analyzed using an Odyssey IR scanner, and signal intensities were quantified using NIH Image/J software.

ELISA assays

To evaluate protein levels of IL-33 in cell lysates and cell culture supernatants, ELISA assays were performed by ELISA kits from R&D Systems according to the manufacturer's instruction.

Flow cytometry analysis

For intracellular cytokine staining, cells were stimulated with PMA (50 ng/ml), lonomycin (250 ng/ml) and Golgi-Plug (1:1000 dilution, BD PharMingen) at 37°C for 4h, followed by fixation with the Fixation/Permeabilization buffer solution (BD Biosciences) according to manufacturer's instruction. The staining procedures for MOG38-49-specific tetramer were previously described⁽²⁾. Stained cells were analyzed on an LSRFortessa (BD Biosciences), and data were analyzed with FlowJo software.

Allogeneic ADSCs treatment and saliva flow rate measurement

Allogeneic ADSCs from mice *C57BL/6J* or *C57BL/6J-II33*^{tm1b(EUCOMM)Cln} treatment and saliva flow rate measurement, For ADSCs treatment, NOD/ShiLtj mice were injected with 1*105 ADSCs (from *C57BL/6* or *C57BL/6J-II33*^{tm1b(EUCOMM)Cln}) in 0.10 ml PBS via the tail vein. Mice were injected with isoproterenol/pilocarpine to stimulate saliva secretion. For saliva flow rate test, mice were weighed, and mild anesthesia was induced with a solution of Ketamine and Xylazine (20 mg/ml; Sigma) in sterile water, given intraperitoneally (1 µl/g of body weight). Salivary secretion was stimulated using 0.1 ml/kg body weight of a pilocarpine solution (50 mg/ml) subcutaneously. Saliva collection began within two minutes of pilocarpine administration. Animals were positioned with a 75-mm hematocrit tube placed in the oral cavity, and the whole saliva was collected into pre-weighed 0.75 ml Eppendorf tubes for 10 minutes. The amount of saliva collected was determined gravimetrically.

Histology

Salivary glands were collected and fixed in 10% formalin (Richard-Allan Scientific), embedded in paraffin, cut longitudinally into 5-mm sections and stained with hematoxylin and eosin. Images were acquired using Aperio digital pathology system and processed in Aperio ImageScope v11.1.2.752 software and the area of inflammatory focus (containing>50 lymphocytes per 4mm² tissue) was calculated per field at X200 magnification by Image-Pro Plus Version 6.0 software (Media Cybernetics). Five entire salivary gland sections for each animal were counted with an average of 10 fields by an expert of histopathology under blinded fashion.

Statistics

All data have a normal distribution and are presented as mean \pm standard error of the mean (SEM; in data of infiltrating area statistics) or standard deviation (in other data) of three independent experiments, and we used an α level of 0.05 for all statistical tests. The mice salivary flow rates were statistically analyzed with repeated measurement; other data were analyzed with student's t-test, one-way analysis of variance (ANOVA), or two-way ANOVA. All *P* value less than 0.05 were considerate significant. Statistical analysis was done with Graphpad Prism 7.

Regent or Resource source identifier

Antibodies Purified anti-mouse CD3 (145-2C11) Bio X Cell Cat# BE0001-1; RRID: AB_1107634 Purified anti-mouse CD28 (37.51) Bio X Cell Cat# BE0015-1; RRID: AB_1107624 Anti-mouse CD4 Alexa Fluor 405(RM4-5) eBioscience Cat# MCD0426; RRID: AB_1473885 Anti-mouse CD4 Alexa Fluor 700 (RM4-5) eBioscience Cat# 56-0042-82; RRID: AB_494000 Anti-mouse CD4 PerCP-Cy5.5 (RM4-5) eBioscience Cat# 45-0042-82; RRID: AB_1107001 Anti-mouse CD4 PE-Cy7 (RM4-5) eBioscience Cat# 25-0042-82; RRID: AB_1107004 Anti-mouse CD8a PerCP-Cy5.5 (53-6.7) eBioscience Cat# 45-0081-82; RRID: AB_1107004 Anti-mouse CD8b FITC (eBioH35-17.2 (H35-17.2)) eBioscience Cat# 11-0083-82; RRID: AB_657764 Anti-mouse CD62L PE (MEL-14) eBioscience Cat# 12-0621-82; RRID: AB_465721 Anti-mouse CD44 APC (IM7) eBioscience Cat# 17-0441-82; RRID: AB_465721 Anti-mouse CD45 Alexa Fluor 700 (30-F11) eBioscience Cat# 56-0451-82; RRID: AB_891454 Anti-mouse CD45 Pacific Blue (30-F11) eBioscience Cat# MCD4528; RRID: AB_1500474 Anti-mouse CD25 PE (PC61.5) eBioscience Cat# 12-0251-82; RRID: AB_465607 Anti-mouse TCRβ APC-Cy7(H57-597) eBioscience Cat# 47-5961-82; RRID: AB_1272173 Anti-mouse TCRγδ Percp-eFlour710 (eBioGL3 (GL-3, GL3)) Cat# 46-5711-82; RRID: AB_2016707 Anti-mouse/rat Foxp3 eFluor 450 (FJK-16a) eBioscience Cat# 48-5773-82; RRID: AB 1518812 Anti-mouse IL-4 PE-Cv7 (11B11) eBioscience Cat# 25-7041-82; RRID: AB 2573520 Anti-mouse IL-10 APC-Cy7 biolegend Cat# 505036; RRID: AB 2566331 Anti-mouse IL-13 PE (eBio13A) eBioscience Cat# 12-7133-82; RRID: AB 763559 Anti-mouse IL-13 PE-eFluor610 (eBio13A) eBioscience Cat# 61-7133-82; RRID: AB 2574654 Anti-mouse IL-17A APC (eBio17B7) eBioscience Cat# 25-7177-82; RRID: AB 10732356 Anti-mouse IFN-y eFluor 450 (XMG1.2) eBioscience Cat# 48-7311-82; RRID: AB 1834366 Anti-mouse TNF-α PE-ef610 (MP6-XT22) eBioscience Cat# 61-7321-82; RRID: AB 2574666 Anti-mouse-Ki-67 PE-ef610 (SoIA15) eBioscience Cat # 61-5698-82; RRID: AB 2574620 Anti-mouse CD3 PerCP-Cyanine5.5 eBioscience Cat # 45-0031-82; RRID: AB 1107000 Anti-mouse Ly-6G/Ly-6C (RB6-8C5), PerCP-Cyanine5.5 eBioscience Cat #45-5931-80; RRID: AB 906247 Anti-mouse CD19 (eBio1D3 (1D3)) PerCP-Cyanine5.5 eBioscience Cat # 45-0193-82; RRID: AB 106999 Anti-mouse B220 (RA3-6B2) PerCP-Cyanine5.5 eBioscience Cat # 45-0452-82; RRID: AB 1107006 Anti-mouse CD11c (N418) PerCP-Cyanine5.5 eBioscience Cat # 45-0114-82; RRID: AB 925727 Anti-mouse CD11b (M1/70) PerCP-Cyanine5.5 eBioscience Cat # 45-0112-82; RRID: AB 953558 Anti-mouse Fc PerCP-Cyanine5.5 biolegend Cat # 149517; RRID: AB 2632750 Anti-mouse TCRβ (H57-597) PerCP-Cyanine5.5 eBioscience Cat # 45-5961-82; RRID: AB 925763 Anti-mouse NK1.1 (PK136) PerCP-Cyanine5.5 eBioscience Cat # 45-5941-82; RRID: AB 914361 Anti-mouse CD127 (A7R34) Alexa Fluor 700 eBioscience Cat # 56-1271-82; RRID: AB 657611 Anti-mouse Ly-6A/E (Sca-1) (D7) FITC eBioscience Cat # 11-5981-82; RRID: AB 465333 Anti-mouse ST2 (RMST2-2) PE eBioscience Cat # 12-9335-82; RRID: AB 2572708 Anti-mouse ST2 (RMST2-2) Percp-eFluor710 eBioscience Cat # 46-9335-82; RRID: AB 2573883 Anti-mouse GATA3 (RMST2-2) Percp-eFluor710 eBioscience Cat # 46-9335-82; RRID: AB 2573883 Anti-IL-33 antibody [Nessy-1] abcam Cat #ab54385; RRID: AB 881109 P44/42 MAPK(Erk1/2) (L34F12) mouse mAb CST Cat# 4696 ; RRID:AB 390780 Phospho- P44/42 MAPK(Erk1/2) (Thr202/Tyr204) (197G2) Rabbit mAb CST Cat# 4377; RRID:AB_331775 p38 MAPK (D13E1) XP Rabbit mAb CST Cat #8690; RRID:AB 10999090 Phospho-p38 MAPK (Thr180/Tyr182) (28B10) Mouse mAb Cat #9216; RRID:AB 331296 Anti-JNK1+JNK2+JNK3 antibody [EPR16797-211] abcam Cat #ab179461; RRID: AB 2744672 Phospho-SAPK/JNK (Thr183/Tyr185) (81E11) Rabbit mAb Cat #4668; RRID:AB 823588 p65 rabbit mAb (C22B4) CST Cat# 4764; RRID:AB 823578 Phospho-p65 (Ser536) rabbit mAb (93H1) CST Cat# 3033; RRID:AB 331284 GAPDH rabbit mAb (14C10) CST Cat# 2118; RRID:AB 561053

Chemicals, Peptides, and Recombinant Proteins

Recombinant human IL-2 R&D Systems Cat# 202-IL-500 Recombinant human TGF-B1 R&D Systems Cat# 240-B-010 Recombinant mouse IL-1ß PeproTech Cat# 211-11B Recombinant mouse IL-4 PeproTech Cat# 214-14 Recombinant mouse IL-5 PeproTech Cat# 215-15 Recombinant mouse IL-6 PeproTech Cat# 216-16 Recombinant mouse IL-7 PeproTech Cat# 217-17 Recombinant mouse IL-12 PeproTech Cat# 210-12 Recombinant mouse IL-13 PeproTech Cat# 210-13 Recombinant mouse IL-15 PeproTech Cat# 210-15 Recombinant mouse IL-21 R&D Systems Cat# 594-ML-025 Recombinant mouse IL-22 R&D Systems Cat# 582-ML-010 Recombinant mouse IL-23 R&D Systems Cat# 1887-ML-010 Recombinant mouse/rat IL-33 R&D Systems Cat# M3300 Recombinant mouse TNF-α PeproTech Cat# 315-01A Recombinant mouse IFN-y PeproTech Cat# 315-05 Recombinant Human IL-1ß PeproTech Cat# 200-01B (5Z)-7-Oxozeaenol R&D Systems Cat# 3604/1 U0126 R&D Systems Cat# 1144/25 SB203580 R&D Systems Cat# 1202/10 SP600125 Invitrogen Cat# tlrl-sp60

JSH23 ab144824 abcam Cat#749886-87-1 PMA Sigma Aldrich Cat# P1585 Ionomycin Sigma Aldrich Cat# I0634 Golgi-Plug BD Biosciences Cat# 555029 TRIzol reagent Invitrogen Cat# 15596026 DNase I Sigma Alcrich Cat# DN25 Collagenase GIBCO Cat# 17104019

Regent or Resource source identifier RNeasy

Mini Kit QIAGEN Cat# 74106 High-Capacity cDNA Reverse Transcription Kit Applied Biosystems Cat# 4368814 TaqMan Gene Expression Master Mix Applied Biosystems Cat# 4369016 Foxp3/Transcription Factor Staining Buffer Set eBioscience Cat# 00-5523-00 Cytofix/Cytoperm Fixation/ Permeabilization Solution Kit BD Biosciences Cat# 554714 CD4⁺ T cell isolation kit, mouse Miltenyi Biotec Cat # 130-104-456 CD4+CD25+ Regulatory T Cell Isolation Kit, mouse Miltenyi Biotec Cat# 130-091-041

Experimental Models: Cell Lines

hADSC Human Adipose-Derived Stem Cells Catalog #: PT-5006, Lonza Walkersville Inc

Experimental Models: Organisms/Strains

Mouse: C57BL/6 The Jackson Laboratory Cat#000664 Mouse: C57BL/6J-II33^{tm1b(EUCOMM)CIn} Washington university at St. Louis Mouse: NOD/ShiLtj The Jackson Laboratory Cat#001976

Oligonucleotides TaqMan

TaqMan Hprt primer, Mm00446968_m1 Applied Biosystems TaqMan IL33 primer Mm00505403_m1 Applied Biosystems TaqMan ST2 primer Mm00516117_m1 Applied Biosystems TaqMan Cebpα, Mm00514283_s1 Applied Biosystems TaqMan Fabp4, Mm00445878_m1 Applied Biosystems TaqMan Ppar, Mm01184322_m1 Applied Biosystems TaqMan Alp, Mm00475834_m1 Applied Biosystems TaqMan Sox9, Mm00448840_m1 Applied Biosystems TaqMan Gapdh, Mm99999915_g1 Applied Biosystems TaqMan Foxp3, Mm00475162_m1 Applied Biosystems TaqMan Human Gapdh Hs02758991_m1 Applied Biosystems TaqMan Human II33, Hs04931857_m1 Applied Biosystems

Regent or Resource source identifier

Software and Algorithms FlowJo 9 software FlowJo, LLC https://www.flowjo.com, RRID:SCR_008520 GraphPad Prism 7 software GraphPad Software https://www.graphpad.com, RRID:SCR_002798 BD LSRFortessa BD Biosciences BD FACSAria cell sorter BD Biosciences

Supplemental References

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