



Article

IL-7 Deficiency Exacerbates Atopic Dermatitis in NC/Nga Mice

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Abstract: Interleukin-7 (IL-7) plays a vital role in the homeostasis of CD4⁺ and CD8⁺ T cells. Although IL-7 has been implicated in T helper (Th)1- and Th17-mediated autoinflammatory diseases, its role in Th2-type allergic disorders, such as atopic dermatitis (AD), remains unclear. Thus, to elucidate the effects of IL-7 deficiency on AD development, we generated IL-7-deficient AD-prone mice by backcrossing IL-7 knockout (KO) B6 mice onto the NC/Nga (NC) mouse strain, a model for human AD. As expected, IL-7 KO NC mice displayed defective development of conventional CD4⁺ and CD8⁺ T cells compared with wild type (WT) NC mice. However, IL-7 KO NC mice presented with enhanced AD clinical scores, IgE hyperproduction, and increased epidermal thickness compared with WT NC mice. Moreover, IL-7 deficiency decreased Th1, Th17, and IFN- γ -producing CD8⁺ T cells but increased Th2 cells in the spleen of NC mice, indicating that a reduced Th1/Th2 ratio correlates with severity of AD pathogenesis. Furthermore, significantly more basophils and mast cells infiltrated the skin lesions of IL-7 KO NC mice. Taken together, our findings suggest that IL-7 could be a useful therapeutic target for treating Th2-mediated skin inflammations, such as AD.

Keywords: IL-7; atopic dermatitis; NC/Nga mice; CD4⁺ T cells; CD8⁺ T cells; IFN- γ ; IL-17



Citation: Park, H.J.; Lee, S.W.; Van Kaer, L.; Lee, M.S.; Hong, S. IL-7 Deficiency Exacerbates Atopic Dermatitis in NC/Nga Mice. *Int. J. Mol. Sci.* **2023**, *24*, 9956. <https://doi.org/10.3390/ijms24129956>

Academic Editors: Kuender D. Yang and Lin-Shien Fu

Received: 25 May 2023

Revised: 6 June 2023

Accepted: 8 June 2023

Published: 9 June 2023



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1. Introduction

Atopic dermatitis (AD) is a Th2-mediated chronic inflammatory skin disease characterized by pruritus, xerosis, erythematosis, and edema. Around 5–15% of all children worldwide, including South Korea, suffer from AD. Apart from genetic factors, such as filaggrin gene mutation, environmental factors (i.e., diet, pollutants, and pathogens) can affect the incidence of AD. Skin barrier function is generally impaired in AD, leading to increased epidermal proliferation and infiltration of Th2-type innate immune cells (i.e., mast cells and basophils). Increased serum IgE levels in AD patients correlate with the severity and relapse of AD. Mast cells and basophils are responsible for eliciting Th2 immune responses in AD due to the high affinity of the IgE receptor (Fc ϵ R1) [1–4]. Allergic immune responses in AD are promoted by Th2 cell-derived IL-4 and IL-5, which can be inhibited by cytokines produced from Th1 and regulatory T (Treg) cells. Thus, the development of AD is closely linked to an imbalance in the ratios of Treg/Th2 cells and Th1/Th2 cells and modulating this balance can affect AD treatment and recovery [5–7].

Interleukin-7 (IL-7) is a critical cytokine in the development and expansion of T cells. It is mainly secreted by thymic and bone marrow stromal cells and plays a significant role in health maintenance and disease prevention [8]. IL-7 signaling deficiency can result in severe immunodeficiency and, following antigen exposure, contributes to memory T cell development and long-term T cell survival [9]. The IL-7 receptor (IL-7R) is a heterodimeric complex consisting of a unique α -chain and the common γ -chain (γ c), which is shared with

the receptors for IL-2, IL-4, IL-7, IL-9, IL-15, and IL-21. While IL-2R is expressed on CD4⁺ T cells, CD8⁺ T cells, and Treg cells, IL-7R is mainly expressed on CD4⁺ T cells and CD8⁺ T cells but not Treg cells [10,11]. In addition, IL-7R signaling can impair the differentiation and suppressive function of Foxp3⁺ Treg cells, and an increase of surface IL-7R expression promotes the skewing of CD4⁺ T cell differentiation into Th17 cells [12].

Previous studies have reported that IL-7R α blockade can mitigate autoimmune diabetes by promoting the proportion of programmed death-1 (PD-1)-expressing effector/memory T cells and Treg cells [13]. In addition, IL-7 signaling is necessary to induce the differentiation of Th1 cells rather than Th17 cells, and serum IL-7 levels correlate inversely with responsiveness to IFN- β therapy in Th1-mediated experimental autoimmune encephalomyelitis (EAE), a mouse model of multiple sclerosis (MS) [14]. Moreover, IL-7 mediates the production of pro-inflammatory cytokines by intra-articular T cells, indicating that IL-7 contributes to increased Th1 responses in patients with rheumatoid arthritis [15].

NC/Nga (NC) mice have been widely employed as a spontaneous murine AD model. When NC mice are maintained under conventional animal housing conditions, they display symptoms similar to AD patients. However, these AD symptoms do not develop under specific pathogen-free (SPF) conditions, suggesting that not only genetic factors but also certain environmental factors (i.e., allergens and bacterial infection) might trigger AD symptoms in NC mice [16].

While most studies on IL-7 have focused on Th1- or Th17-mediated autoinflammatory diseases, its role in Th2-mediated diseases, such as AD, is not fully understood. Therefore, in this study, we investigated the immunoregulatory effects of IL-7 on effector T cells in AD development.

2. Results

2.1. IL-7 Deficiency Exacerbates AD Pathogenesis in NC/Nga Mice

A previous study demonstrated that IL-7 signaling blockade using anti-IL-7R α monoclonal antibody (mAb) treatment or endothelial-specific deletion of IL-7R α attenuates psoriasis-like skin inflammation mediated by infiltration of T cells and DCs into the inflamed lesion [17]. Based on these promising findings, we investigated whether IL-7 plays a role in another skin inflammatory disease, AD. To address this issue, we generated IL-7 knockout (KO) NC mice by backcrossing the mutant IL-7 allele from mice on a B6 genetic background onto the AD-susceptible NC genetic background. Both wild-type (WT) and IL-7 KO NC mice developed skin inflammation spontaneously when raised under conventional housing conditions for eight weeks (Figure 1A). While being maintained under conventional housing conditions, these mice were monitored to measure the clinical skin score, including lichenification, edema, erosion/excoriation, scarring/dryness, and erythema/hemorrhage, every two weeks from six to fourteen weeks of age. While AD severity in WT NC mice increased moderately, IL-7 KO NC mice exhibited a more severe form of AD with increased skin inflammation (Figure 1B). Moreover, IL-7 KO NC mice displayed increased clinical disease scores and serum IgE levels compared with WT NC mice (Figure 1C,D). In addition, we measured the epidermal thickness in hematoxylin and eosin (H&E)-stained skin sections of both WT and IL-7 KO NC mice with AD. The skin lesions of IL-7 KO NC mice displayed a significant increase in epidermal thickness compared with those of WT NC mice (Figure 1E,F). WT NC mice housed under SPF conditions were used as a negative control for AD development. As expected, no clinical AD symptoms were detected in SPF NC mice (Figure 1B,D). Our findings suggest that IL-7 deficiency contributes to the modulation of AD development based on our clinical observations of skin inflammation and serum IgE levels.

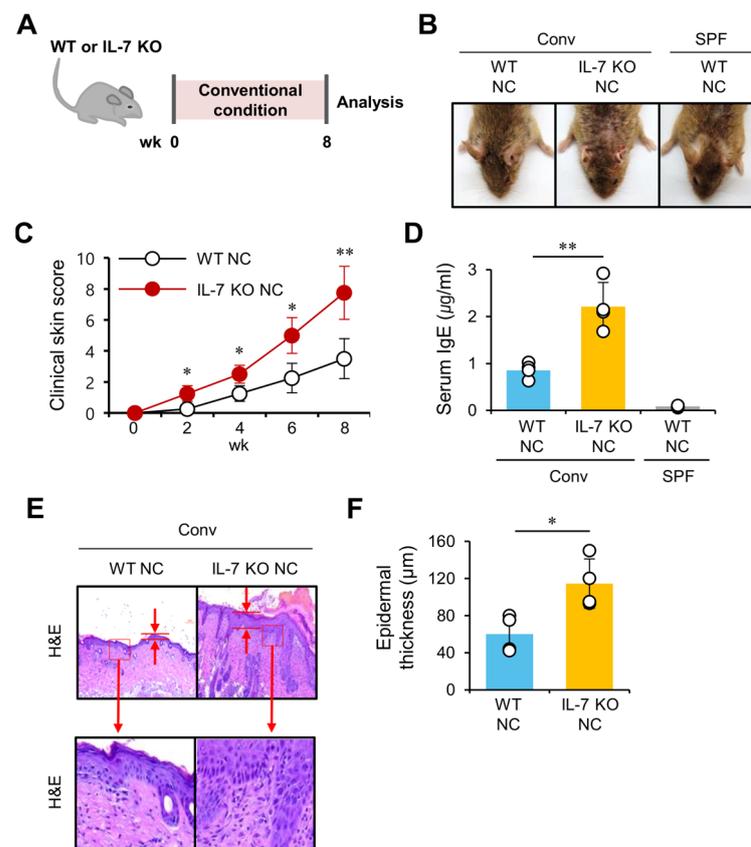


Figure 1. Exacerbation of AD development in IL-7 KO NC mice. (A) Experimental outline. WT or IL-7 KO NC mice were allowed to develop AD spontaneously under conventional housing conditions as described in Materials and Methods. WT NC mice housed under specific pathogen-free (SPF) conditions were used as a negative control for AD development. (B,C) The clinical symptoms were measured every two weeks to monitor the onset of AD. (D) Serum IgE levels were measured using ELISA. (E,F) Skin samples were prepared from WT NC or IL-7 KO NC mice. (E) The skin lesions were sectioned and stained with H&E. (F) The epidermal thickness was measured in 10 random high-power fields ($400\times$) per sampled lesion using Image J software (version 1.53s, imagej.nih.gov/ij/; accessed on 16 June 2022). The mean values \pm SD ($n = 4$ in A–F; per group in the experiment; Student’s *t*-test; * $p < 0.05$, ** $p < 0.01$) are shown. One representative experiment of two experiments is shown.

2.2. IL-7 Deficiency Induces Impaired Development of CD4⁺ and CD8⁺ T Cells in NC Mice with AD

As it has been reported that IL-7 signaling can enhance the survival of T cells by up-regulating anti-apoptotic Bcl-2 and down-regulating pro-apoptotic Bim [18], we examined whether IL-7 deficiency affects T cell profiles (phenotypes) in NC mice. For this purpose, total T cells were isolated from NC mice with AD and assessed for CD4 and CD8 expression using flow cytometry. We found that IL-7 KO NC mice displayed smaller spleen size and decreased frequency of splenocytes (Figure 2A) when they developed AD. Furthermore, we found that the number and frequency of both CD4⁺ T cells and CD8⁺ T cells were significantly lower in IL-7 KO NC mice than in WT NC mice during AD development (Figure 2B,C). To evaluate the effect of AD development on impaired T cell development mediated by IL-7 deficiency, we compared the extent of T cell development in WT NC and IL-7 KO NC housed under SPF conditions. We found that IL-7 deficiency-mediated impairment of T cell development was not affected by AD development (Figure S1). Interestingly, however, both WT NC and IL-7 KO NC mice with AD displayed reduced CD4⁺ and CD8⁺ T cell numbers compared with their respective controls without AD. Consistent with previous studies, our results indicate that IL-7 is essential in maintaining T cells in

NC mice under both SPF and conventional housing conditions. Thus, it appears that exacerbated AD pathogenesis in IL-7 KO NC mice is not attributed to a simple quantitative reduction of T cells but rather to qualitative changes of T cell pools.

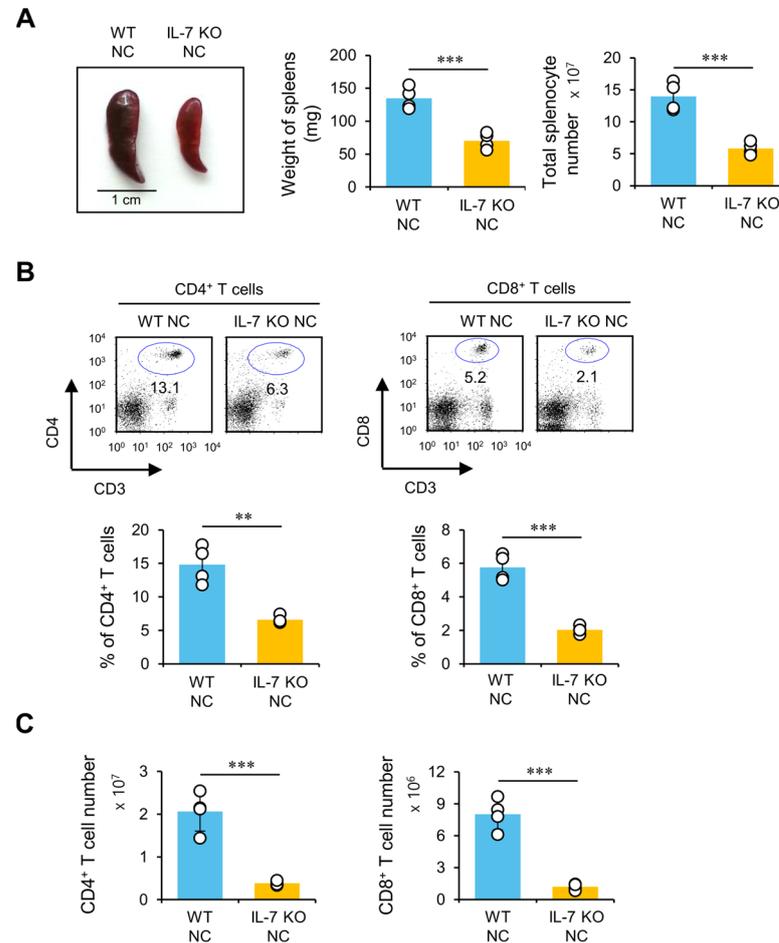


Figure 2. Impaired T cell development in IL-7 KO NC mice with AD. (A–C) WT NC and IL-7 KO NC mice were transferred to the conventional housing conditions at six weeks of age to develop AD spontaneously. Splenocytes were prepared from these mice at 14 weeks of age. (A) (Left), A representative picture of the spleens from AD-induced WT NC and IL-7 KO NC mice. (Middle) and (Right), Spleen weights and splenocyte numbers of these mice. (B,C) The percentages and the absolute total cell numbers of both CD4⁺ and CD8⁺ T cells were analyzed via flow cytometry. The mean values \pm SD ($n = 4$ in A–C; per group in the experiment; Student's *t*-test; ** $p < 0.01$, *** $p < 0.001$) are shown. One representative experiment of two experiments is shown.

2.3. Worsened AD Pathogenesis in IL-7 KO NC Mice Is Associated with Th2-Biased Immune Responses and Down-Regulation of IFN- γ and IL-17 Expression

As previous studies have shown that elevated Th2 immune responses are responsible for AD pathogenesis [5], we investigated whether IL-7 deficiency impacts Th2 cell differentiation during AD development. To explore this possibility, we performed flow cytometric analyses to determine Th1/Th2 cytokine profiles of CD4⁺ T cells from IL-7 KO NC mice with fully developed AD. We found that IL-7 KO NC mice with AD exhibit reduced production of IFN- γ and IL-17 cytokines in CD4⁺ T cells compared to WT NC mice, suggesting that Th1 and Th17 cell differentiation are negatively affected in IL-7 KO NC mice during AD development (Figure 3A). Next, we also measured Th2 cytokines (IL-4 and IL-5) in CD4⁺ T cells to examine whether exacerbated AD in IL-7 KO NC mice is related to changes in Th2 cells. We found that IL-7 KO NC mice display elevated IL-4 and IL-5 expression in CD4⁺ T cells following AD induction compared to WT NC mice

(Figure 3B). IL-7 KO NC mice displayed a reduced Th1/Th2 ratio, indicating that Th2-biased immune responses dominate in these mice (Figure 3C). Interestingly, we also found decreased IFN- γ -producing but not IL-17-producing CD8⁺ T cells in the IL-7 KO NC mice, similar to the observed reduction in Th1 cells (Figure 3D). In addition, to evaluate the effect of AD susceptibility (but not active AD) on T cell cytokine production, we compared cytokine production of CD4⁺ and CD8⁺ T cells in WT NC and IL-7 KO NC mice housed under SPF conditions. We found that IL-7 KO NC without AD display a reduction in IFN- γ and IL-17 production by CD4⁺ T cells. However, unlike WT NC and IL-7 KO NC mice with AD, the control NC mice housed under SPF conditions did not show any significant changes in IL-4 and IL-5 expression by CD4⁺ T cells, which indicates that exacerbation of AD might be attributed to environmental factors, such as microbial infections (Figure S2). These results indicate that IL-7 deficiency up-regulates pathogenic Th2 immune responses but down-regulates Th1 and CD8⁺ T cells in NC mice during AD development.

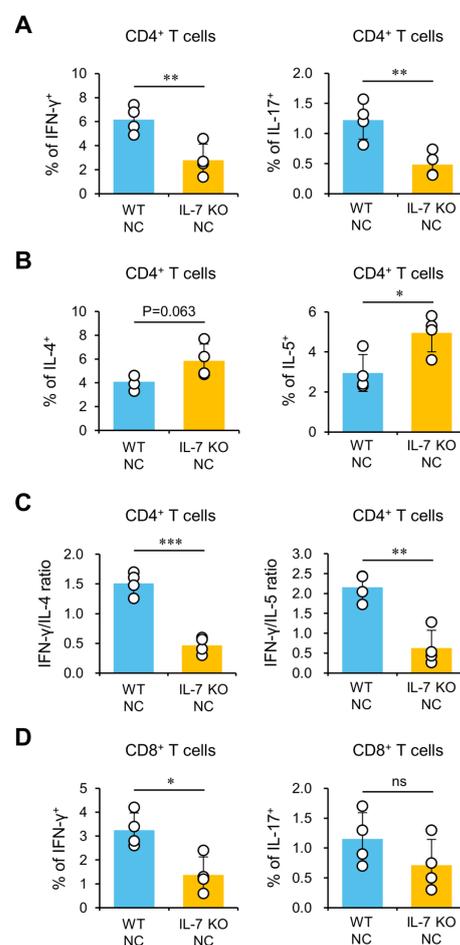


Figure 3. Down-regulation of IFN- γ and IL-17 expression may be linked to up-regulation of Th2-biased immune responses in IL-7 KO NC mice. (A–D) WT NC and IL-7 KO NC mice were transferred to the conventional housing conditions at six weeks of age to develop AD spontaneously. Splenocytes were prepared from these mice at 14 weeks of age. (A) IFN- γ - and IL-17-producing subpopulations among splenic CD4⁺ T cells from each group were determined using flow cytometry. (B) IL-4- and IL-5-producing subpopulations among splenic CD4⁺ T cells were evaluated using flow cytometry. (C) The ratio of the IL-4- or IL-5-producing population to the IFN- γ -producing population in splenic CD4⁺ T cells was measured using flow cytometric analysis. (D) IFN- γ - and IL-17-producing subpopulations among splenic CD8⁺ T cells from each group were determined using flow cytometric analysis. The mean values \pm SD ($n = 4$ in A–D; per group in the experiment; Student's *t*-test; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$) are shown. One representative experiment of two experiments is shown. ns, not significant.

2.4. The Effects of IL-7 Deficiency on AD Severity Are Associated with Expanded Basophils and Mast Cells in the Skin

Basophils and mast cells can contribute to allergic immune responses through IgE-mediated FcεR1 stimulation. Moreover, basophils can enhance Th2 polarization by quickly secreting IL-4. Thus, we investigated whether these innate immune cells correlate with exacerbated AD pathogenesis in IL-7 KO NC mice. As expected, splenic basophils and mast cells were significantly increased in number in IL-7 KO NC mice than in WT NC mice during AD development (Figure 4A). Additionally, IL-7 KO NC mice had a more significant infiltration of basophils and mast cells in skin lesions (Figure 4B). These findings indicate that IL-7 deficiency promotes Th2-mediated immune responses and consequently expands allergic effectors, such as basophils and mast cells, ultimately exacerbating spontaneous AD development in IL-7 KO NC mice.

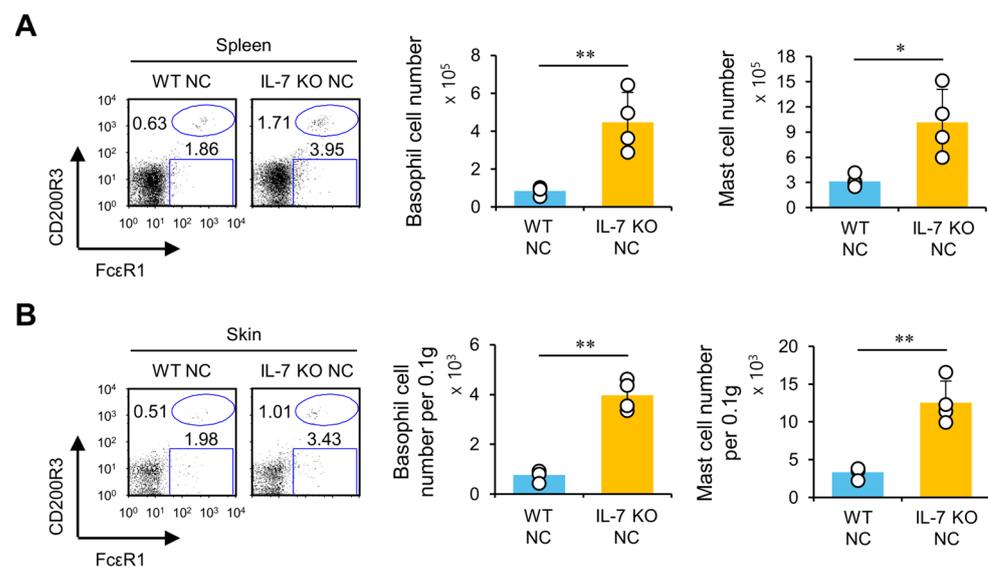


Figure 4. IL-7 deficiency is associated with expanded basophils and mast cells in NC mice with AD. **(A)** Spleens were prepared from mice as shown in Figure 2. The frequency and total absolute cell number of splenic basophils (FcεR1⁺CD200R3⁺CD3⁻CD19⁻) and mast cells (FcεR1⁺CD200R3⁻CD3⁻CD19⁻) were determined using flow cytometry. **(B)** Mononuclear cells (MNCs) in the skin were isolated from mice as shown in Figure 1. The absolute cell numbers of basophils (CD45⁺FcεR1⁺CD200R3⁺CD3⁻CD19⁻) and mast cells (CD45⁺FcεR1⁺CD200R3⁻CD3⁻CD19⁻) were determined in the skin using flow cytometry. The mean values ± SD ($n = 4$ in A,B; per group in the experiment; Student's *t*-test; * $p < 0.05$, ** $p < 0.01$) are shown. One representative experiment of two experiments is shown.

2.5. Human Gene Expression Data Suggest a Strong Association between IL-7 Gene Expression and IFN-γ and IL-17 Gene Expression

As our results strongly suggest that IL-7 plays a critical role in determining the effector functions of CD4 and CD8 T cells, we next asked whether IL-7 gene expression might be linked to Th1/Th17 (IFN-γ and IL-17) and Th2 (IL-4 and IL-5) cytokine expression profiles in humans. To examine a possible correlation between IL-7 gene expression and T helper cytokine gene expression (*IFNG*, *IL-17*, *IL-4*, and *IL-5*), we took advantage of the data from the GEPIA (Gene Expression Profiling Interactive Analysis) website containing expression data for human spleen. Our analysis revealed that *IL-7* expression considerably correlated with the expression of *IFNG* ($R = 0.53$, $p = 1.3 \times 10^{-8}$) and *IL-17A* ($R = 0.21$, $p = 0.032$), but not with *IL-4* and *IL-5* (Figure 5A,B). These results support the notion that IL-7 regulates T helper cell differentiation, ultimately contributing to the outcome of AD pathogenesis.

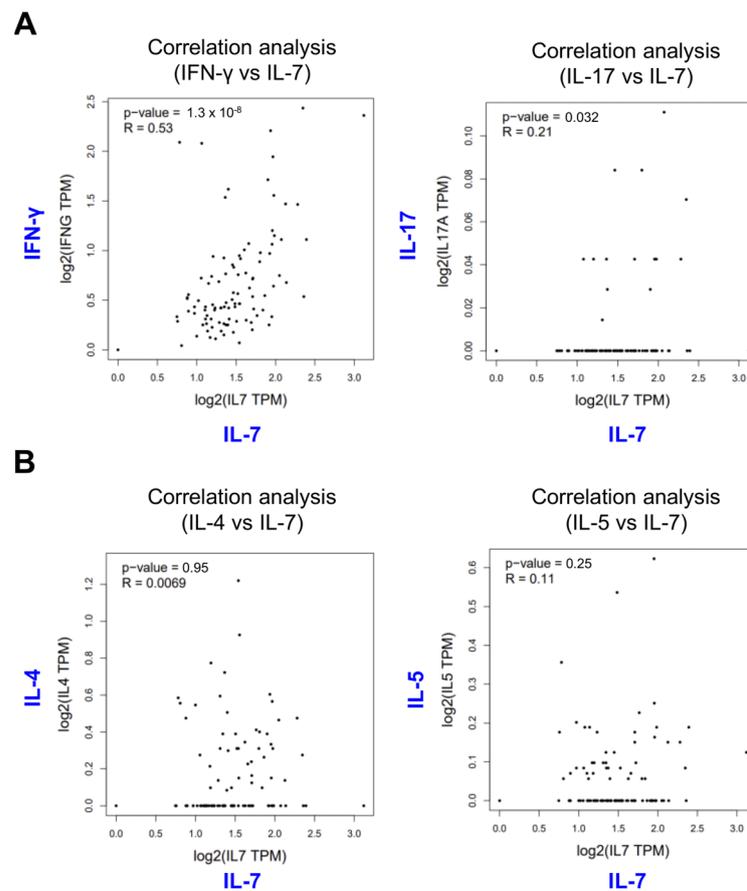


Figure 5. Correlation analyses of *IL-7* and *IFNG* or *IL-17A* gene expression in human spleen. (**A,B**) Pearson correlation analysis of *IL-7* and Th-related cytokine genes (*IFNG*, *IL-17A*, *IL-4*, and *IL-5*) was conducted using the human splenic data from the GEPIA (<http://gepia.cancer-pku.cn/index.html>, accessed on 9 August 2022) tool (TPM; transcripts per million reads).

3. Discussion

In this study, we investigated the role of *IL-7* on AD development by employing *IL-7* KO NC mice. NC mice show an impaired immune response against bacterial skin infections, such as *Staphylococcus aureus* (the leading cause of AD in these mice) [19]. Protective immune responses against invasive *S. aureus* infection require *IFN- γ* and *IL-17* [20,21], which act in synergy to safeguard the oral mucosal layer against *S. aureus* [22]. Our results indicate that the reduction of *IFN- γ* - and *IL-17*-producing $CD4^+$ T cells in *IL-7* KO NC mice may contribute to the breakdown of the skin's immune defense system against *S. aureus* colonization. Thus, it would be interesting to investigate whether the severe skin inflammation found in *IL-7* KO NC mice is attributed to *S. aureus*-dominant colonization in the skin.

Furthermore, as splenic $CD4^+$ T cells from *IL-7* KO mice are defective in producing *IFN- γ* and *IL-17* rather than Th2-type cytokines (e.g., *IL-4* or *IL-5*), the decrease of *IFN- γ* and *IL-17* production may trigger the initiation of Th2 immune responses, consequently leading to accelerated AD development in *IL-7* KO mice (Figure S2). Moreover, consistent with our results, GEPIA analysis of human spleen showed *IL-7* production significantly correlates with *IFN- γ* or *IL-17* but not with *IL-4* or *IL-5* (Figure 4). These findings suggest that a systemic deficiency of *IL-7* down-regulates *IFN- γ* and *IL-17* production by $CD4^+$ T cells, likely providing a Th2-conducive environment that might be favorable for AD-exacerbating invasive *S. aureus* infection.

Systemic oral immunosuppressive drugs are commonly used to treat AD, but more than 70% of AD patients receiving these therapies exhibit lymphopenia, a quantitative

decrease in lymphocyte cell numbers ($\leq 1 \times 10^3$ cells/ μ L) [23]. Recent findings also demonstrated that patients with atopic eczema (AE) have lower lymphocyte counts than individuals without AE, suggesting that a decrease in lymphocyte cell numbers significantly correlates with more severe AD [24]. It has been reported that IL-7 therapy can increase functional T cells and reverse profound lymphopenia [25,26]. Moreover, lymphopenia in AD patients shows a tendency for preferential depletion of IFN- γ -producing T cells (both CD4⁺ and CD8⁺ T cells), which might result in Th2-deviated immune responses [27]. Based on the capacity of IFN- γ to negatively regulate both the activation and survival of AD-initiating effector cells (e.g., mast cells and basophils) [28,29], the IL-7 deficiency-mediated reduction in IFN- γ -producing T cells might be responsible for mast cell and basophil activation, which consequently could enhance AD development. As our results indicate that AD development in IL-7 KO NC mice is accompanied by a decreased number of CD4⁺ and CD8⁺ T cells, it would be worthwhile to investigate whether IL-7 is related to the onset of lymphopenia in AD patients.

Intriguingly, a recent study reported that IL-7 is critical for maintaining invariant natural killer T (iNKT) cells, a subset of glycolipid-reactive T cells with innate-like functions, in peripheral tissues, including lymph nodes, spleen, liver, and lung [30]. Previous studies have revealed that iNKT cells are impaired in NC mice due to the absence of the TCR V β 8 gene [31]. Recently, our studies have shown that V α 14 TCR transgenic (V α 14^{Tg}) NC mice overexpressing V α 14 iNKT cells exhibit significantly reduced skin inflammation upon spontaneous AD development compared to WT NC mice [3,32,33]. Furthermore, skin iNKT cells from V α 14^{Tg} NC mice exhibit a Th1-like phenotype with high IFN- γ and IL-2 during AD development [32]. Thus, it might be interesting to examine whether IL-7 deficiency can trigger the protective effects of V α 14^{Tg} iNKT cells on AD development in NC mice.

Our findings indicate that IL-7 KO NC mice develop worsened skin inflammation and a faster disease progression. We propose that the deficiency of IL-7 expression contributes to the development of AD through a series of steps. Firstly, a decrease in IFN- γ - and IL-17-producing CD4⁺ T cells occurs due to reduced IL-7 expression, possibly resulting in uncontrolled *S. aureus* infection. Secondly, IL-4-producing basophils and mast cells expand, which promotes the differentiation of naive CD4⁺ T cells into Th2 cells. Finally, impaired IFN- γ production induced by IL-7 deficiency causes a loss of negative regulation of IL-4 activity, leading to uncontrollable AD development. Consequently, our research demonstrates that IL-7 has the potential to be used as a therapeutic approach for managing inflammatory skin diseases, such as AD. Thus, IL-7 immunotherapy might be an option to substitute steroid therapy, which is being widely used despite its adverse effects.

4. Materials and Methods

4.1. Study Design

This study was designed to investigate the effect of IL-7 on the development of AD in IL-7 KO NC mice. To address this issue, AD was allowed to develop spontaneously in IL-7 KO mice and WT controls. Skin leukocytes and splenocytes were then isolated and analyzed using flow cytometry, while sera were harvested and analyzed using ELISA. This study received approval from the Sejong University Institutional Review Board before the experiments were conducted (SJ-20181101E2).

4.2. Mice

WT NC mice were purchased from Jung Ang Lab Animal Inc. (Seoul, Republic of Korea). IL-7 KO B6 mice [34] were backcrossed to NC mice for more than ten generations. These mice were maintained at the Sejong University and were used for experiments at 6–12 weeks of age. The mice were maintained on a 12 h light/12 h dark cycle in a temperature-controlled barrier facility with free access to food and water. They were fed a γ -irradiated sterile diet and provided with autoclaved tap water. To monitor spontaneous AD development, mice were transferred to a conventional animal facility at six weeks of age. Age- and sex-matched mice were used for all experiments. The animal experiments

were approved by the Institutional Animal Care and Use Committee of Sejong University (SJ-20181101E2).

4.3. Flow Cytometry

The following mAbs were obtained from BD Biosciences (San Jose, CA, USA): Phycoerythrin (PE)-, PE-Cy7- or allophycocyanin (APC)-conjugated anti-CD3 ϵ (clone 145-2C11); PE-Cy7- or APC-conjugated anti-CD4 (clone RM4-5); PE-Cy7- or APC-conjugated anti-CD8 α (clone 53-6.7); PE-Cy7- or APC-conjugated anti-CD45 (clone 30-F11), PE-conjugated immunoglobulin G (IgG) (isotype control) (clone R3-34). In addition, the following mAbs from Thermo Fisher Scientific (Waltham, MA, USA) were used: APC-conjugated anti-CD200R3 (clone Ba13); PE-Cy7-conjugated anti-CD19 (clone ID3); PE-conjugated anti-Fc ϵ R1 (clone MAR-1); PE-conjugated anti-IFN- γ (clone XMG1.2); PE-conjugated anti-IL-4 (clone BVD6-24G2); PE-conjugated anti-IL-5 (clone TRFK5); PE-conjugated anti-IL-17 (clone eBio17B7). To perform surface staining, cells were harvested and washed twice with cold 0.5% BSA-containing PBS (FACS buffer). To block the Fc receptor, the cells were incubated with anti-CD16/CD32 mAbs on ice for 10 min and subsequently stained with fluorescently labeled mAbs. Flow cytometric data were acquired using a FACSCalibur flow cytometer (Becton Dickinson, San Jose, CA, USA) and analyzed using FlowJo software (version 8.7; Tree Star, Ashland, OR, USA) [35].

4.4. Flow Cytometry

For intracellular staining, splenocytes were incubated with brefeldin A, an intracellular protein transport inhibitor (10 μ g/mL), in RPMI medium (Gibco BRL, Gaithersburg, MD, USA) for 2 h at 37 $^{\circ}$ C. The cells were stained for cell surface markers, fixed with 1% paraformaldehyde, washed once with cold FACS buffer, and permeabilized with 0.5% saponin. The permeabilized cells were then stained for an additional 30 min at room temperature with the indicated mAbs (PE-conjugated anti-IFN- γ , anti-IL-4, anti-IL-5, anti-IL-17, or PE-conjugated isotype control rat IgG) [36]. More than 5000 cells per sample were acquired using a FACSCalibur flow cytometer, and the data were analyzed using the FlowJo software package (version 8.7; Tree Star, Ashland, OR, USA).

4.5. ELISA

Serum IgE levels were measured with a sandwich ELISA (clone R35-72 for capturing IgE and R35-118 for detecting IgE; BD PharMingen, San Jose, CA, USA). In addition, the optical density was measured at 450 nm with an immunoreader (Bio-Tek ELX-800, Winooski, VT, USA).

4.6. Preparation of Skin Cell Suspensions

The skin was dissected and dermal fat was removed with scissors. The tissue was cut into small pieces with a scalpel and digested with 2.5 mg/mL collagenase type IV (Sigma, St. Louis, MO, USA) and 1 mg/mL DNase I (Promega, Madison, WI, USA) for 4 h at 37 $^{\circ}$ C. At the end of the incubation, the digested tissue was dissociated into single-cell suspensions using gentleMACS Dissociator (Miltenyi, Bergisch Gladbach, Germany) in combination with C Tubes. Single-cell suspensions were smashed through a 70 μ m nylon cell strainer (BD Falcon, Franklin Lakes, NJ, USA) and collected in a 50 mL Falcon tube. The cells were washed once with PBS plus 10 % FBS (1400 rpm, 10 min, 4 $^{\circ}$ C) and subsequently separated with 37%/70% Percoll (GE Healthcare, Chicago, IL, USA) gradients. Mononuclear cells were collected from the layer below 37% and above 70% gradient. After washing with PBS, the total mononuclear cell number was determined using a hemacytometer with 0.4% trypan blue (Welgene, Gyeongsan-si, Republic of Korea) before antibody staining.

4.7. Analysis of Skin Sections

The dorsal skin was fixed in 4% paraformaldehyde, embedded in paraffin, and cut into six μ m sections using a microtome (RM 2235, Leica, Wetzlar, Germany). The sections

were then stained with hematoxylin and eosin (H&E) to analyze histological changes. The epidermal thickness was measured in ten high-power fields (400×) per each section using Image J software (version 1.53s, imagej.nih.gov/ij/; accessed on 16 June 2022) (National Institutes of Health, Bethesda, MD, USA).

4.8. Scoring the Severity of Skin Lesions

Skin lesions were scored at the indicated time points. The scoring was based on the severity of lichenification, edema, erosion/excoriation, scarring/dryness, and erythema/hemorrhage. The total clinical skin severity score was defined as the sum of the five signs (none = 0; mild = 1; moderate = 2; and severe = 3).

4.9. Data Collection in the GEPIA (Gene Expression Profiling Interactive Analysis)

Correlation analysis between *IL-7* and Th-related cytokines (*IFNG*, *IL-17A*, *IL-4*, and *IL-5*) was conducted using RNA-seq data of healthy human spleen from a bioinformatics GEPIA database [37]. Data credit: GEPIA. Data summary images were obtained from <http://gepia.cancer-pku.cn/index.html> (accessed on 9 August 2022).

4.10. Statistical Analysis

Statistical significance was determined using Excel (Microsoft, Redmond, WA, USA). Student's *t*-test was performed for the comparison of two groups (* $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$ were considered significant in the Student's *t*-test). Two-way ANOVA analysis was carried out using the VassarStats (<http://vassarstats.net/anova2u.html>). # $p < 0.05$, ## $p < 0.01$, and ### $p < 0.001$ were considered to be significant in the two-way ANOVA.

Supplementary Materials: The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/ijms24129956/s1>.

Author Contributions: Conceptualization, H.J.P., S.W.L. and S.H.; methodology, H.J.P., S.W.L., M.S.L. and S.H.; software, H.J.P., S.W.L. and S.H.; validation, H.J.P., S.W.L. and S.H.; formal analysis, H.J.P., S.W.L. and S.H.; investigation, H.J.P., S.W.L., L.V.K., M.S.L. and S.H.; resources, H.J.P., S.W.L. and S.H.; data curation, H.J.P., S.W.L. and S.H.; writing—original draft preparation, H.J.P., S.W.L. and S.H.; writing—review and editing, H.J.P., S.W.L., L.V.K., M.S.L. and S.H.; visualization, H.J.P., S.W.L. and S.H.; supervision, S.H.; project administration, H.J.P., S.W.L. and S.H.; funding acquisition, H.J.P., S.W.L. and S.H. All authors have read and agreed to the published version of the manuscript.

Funding: This work was supported by the Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education (NRF-2021R1I1A1A01054418 to S.W.L.; NRF-2021R1I1A1A01051465 to H.J.P.; NRF-2019R1A2C1009926 and NRF-2022R1A2C1009590 to S.H.).

Institutional Review Board Statement: The study was conducted according to the guidelines on the care and use of laboratory animals approved by the Institutional Animal Care and Use Committee of Sejong University (approved on 1 November 2018).

Informed Consent Statement: Not applicable.

Data Availability Statement: The data will be available from the corresponding author upon reasonable request.

Conflicts of Interest: L.V.K. is a member of the scientific advisory board of Isu Abxis Co., Ltd. (Republic of Korea). The other authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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