Cornea

TSLP Protects Corneas From Pseudomonas aeruginosa Infection by Regulating Dendritic Cells and IL-23-IL-17 **Pathway**

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PURPOSE. We sought to determine the role of epithelium-produced thymic stromal lymphopoietin (TSLP) and its underlying mechanisms in corneal innate immune defense against Pseudomonas (P.) aeruginosa keratitis.

METHODS. The expression of TSLP and TSLPR in cultured human corneal epithelial cells (HCECs) and mouse corneas was determined by PCR, Western, and/or ELISA. Cellular localization of TSLP receptor (TSLPR) was determined by whole mount confocal microscopy. TSLP-TSLPR signaling was downregulated by neutralizing antibodies and/or small interfering (si)RNA; their effects on the severity of *P. aeruginosa*-keratitis and cytokine expression were assessed using clinical scoring, bacterial counting, PMN infiltration, and real-time PCR. The role of dendritic cells (DCs) in corneal innate immunity was determined by local DC depletion using CD11c-DTR mice.

Results. P. aeruginosa-infection induced the expression of TSLP and TSLPR in both cultured primary HCECs and in C57BL/6 mouse corneas. While TSLP was mostly expressed by epithelial cells, CD11c-positive cells were positive for TSLPR. Targeting TSLP or TSLPR with neutralizing antibodies or TSLPR with siRNA resulted in more severe keratitis, attributable to an increase in bacterial burden and PMN infiltration. TSLPR neutralization significantly suppressed infection-induced TSLP and interleukin (IL)-17C expression and augmented the expression of IL-23 and IL-17A. Local depletion of DCs markedly increased the severity of keratitis and exhibited no effects on TSLP and IL-23 expression while suppressing IL-17A and C expression in P. aeruginosa-infected corneas.

CONCLUSIONS. The epithelium-expressed TSLP plays a protective role in *P. aeruginosa* keratitis through targeting of DCs and in an IL-23/IL-17 signaling pathway-related manner.

Keywords: bacterial keratitis, TSLP, dendritic cells, innate immunity, IL-23-IL-17 pathway

 $B_{a}^{\rm acterial\ keratitis\ is\ a\ common\ infectious\ ocular\ disease\ and\ a\ leading\ cause\ of\ ocular\ morbidity\ and\ blindness$ worldwide.¹ The Gram-negative pathogen, *P. aeruginosa* is a major corneal pathogen, representing nearly 38% of infectious keratitis cases.¹ P. aeruginosa keratitis is generally thought to be more severe at presentation, more difficult to treat, and results in worse visual outcomes than other forms of bacterial keratitis.² Although it is a ubiquitous pathogenic bacterium, P. aeruginosa can only invade the cornea to cause opportunistic infection when the corneal epithelium barrier is impaired. Like other mucosal linings, corneal epithelial cells express pathogen recognizing receptors and play an important role in the innate immune response by sensing the presence of pathogens and by expressing inflammatory cytokines such as IL-1 β , IL-8 and CXCL2, and antimicrobial peptides (AMPs) such as β -defensins and CRAMP.³ In addition to epithelial cells, the epithelial layer contains a rich network of sensory nerves and sparselydistributed intraepithelial dendritic cells. These cells form a functional unit with a coordinated response to environmental challenges, including infectious pathogens. The molecules

involved in the coordination of this defense network remain largely to be elucidated.

TSLP is an IL-7 like cytokine expressed mainly by epithelial cells and is known to activate STAT3, STAT5, and JAK2 pathways, which control processes such as cell proliferation and development of the hematopoietic system.⁴ Originally shown to promote the growth and activation of B cells, it is now known to have wide-ranging impacts on both hematopoietic and nonhematopoietic cell lineages, including dendritic cells, basophils, eosinophils, mast cells, CD4(+), CD8(+) and natural killer T cells, B cells and epithelial cells.⁴ TSLP-induced Th2 responses are associated with the pathogenesis of allergic inflammatory diseases, including atopic dermatitis, asthma, and rhinitis. It can directly or indirectly promote Th2 and Treg responses, and inhibit Th1 and Th17 responses through limiting the expression of pro-inflammatory cytokines such as IL-17 and IFN-gamma.⁵ Based on recent findings in humans and mouse models, TSLP might also be involved in the pathogenesis of inflammatory bowel disease and progression of cancer.⁶ The receptor of TSLP, mainly found in dendritic cells, is a heterodimeric complex that consists of TSLP receptor (TSLPR)

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and IL-7R α . Activation of this complex results in STAT3 and STAT5 activation.^{7,8} Evidence shows that epithelial cells increasingly express TSLP mRNA and protein upon stimulation by microbial products.⁹ For example, the colon of TSLPR^{-/-} mice displays exaggerated Th1/Th17 responses and reduced Treg cell activation even in the presence of a limited and benign bacterial community.¹⁰ In addition, an alternatively spliced short form of human TSLP has been shown to act as a potent antimicrobial peptide.^{11,12} Expression of TSLP in the skin, oral and GI mucosa is part of the defense barrier that aids in the control of both commensal and pathogenic microbes.¹¹ In the cornea, epithelium-derived TSLP has been linked to experimental mouse allergic conjunctivitis.^{13,14} and *Aspergillus fumigatus* keratitis in in vitro cell culture models.^{15,16} The involvement of TSLP in bacterial keratitis in vivo remains unknown.

We addressed here whether TSLP has a role in regulating *P. aeruginosa* keratitis. In this study, we investigated the expression of TSLP and TSLPR in C57BL/6 mouse corneas in response to *P. aeruginosa* infection and demonstrated that targeting the TSLP-TSLPR signaling axis significantly increased the severity of keratitis including leading to a high bacterial burden. Our data suggests that TSLP plays a protective role in the cornea from *P. aeruginosa* infection through activation of DCs.

MATERIALS AND METHODS

Animals

Wild-type C57BL/6 mice (8 weeks; female) were purchased from the Jackson Laboratory (Bar Harbor, ME, USA). B6-DTR mice, which express simian diphtheria toxin receptor (*DTR*)enhanced green fluorescent protein fusion protein under the control of the *CD11c* promoter, were originally purchased from the Jackson Laboratory and breaded in house at a Wayne State University animal housing facility. All animal procedures were performed in compliance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research and were approved by the Institutional Animal Care and Use Committee of Wayne State University.

Bacterial Preparation

P. aeruginosa strain ATCC 19660 (cytotoxic) was used in this study. This strain is a standard laboratory strain that provides a reproducible inflammatory response in the cornea in the B6 mouse.¹⁷⁻¹⁹ A colony of bacteria was picked and grown overnight at 37°C in tryptic soy broth (TSB) with constantly shaking. The next morning, 20 μ L of *P. aeruginosa*-containing TSB was transferred to 4 mL fresh medium and allowed to grow for 4 hours with OD₆₀₀ less than 1. After the concentration of bacteria in TSB was determined (1 OD₆₀₀ = \times 10⁸ *P. aeruginosa*/mL), the bacteria were washed and resuspended in PBS and then either used to infect mouse corneas or placed in a boiling water bath for 5 minutes to generate heat-killed *P. aeruginosa*. Heat-killed bacteria were diluted with growth factor-free and antibiotic-free keratinocyte basic medium (KBM; Lonza, Basel, Switzerland).

Cell Culture

Primary human corneal epithelial cells (HCECs) were isolated from human donor corneas obtained from Eversight Michigan (Ann Arbor, MI, USA) using a previously described method.²⁰ P4 of the primary HCECs were used for the experiments. Before treatment, cells were starved overnight in KBM. Subsequently, cells were challenged with heat-killed *P. aeruginosa* (1:50 multiplicity of infection) for the indicated times.

Mouse Model of P. aeruginosa Keratitis

Mice, five in each group, were anesthetized with an intraperitoneal injection of ketamine (90 mg/kg) and xylazine (10 mg/kg) before surgical procedures. Mouse corneas were scratched gently with a sterile 26-gauge needle to create three 1-mm incisions to break the epithelial barrier and inoculated with 1.0×10^4 colony-forming units (CFUs) of *P. aeruginosa* in 5 µL PBS. The number of bacteria used to inoculate the mouse corneas were 100-fold less than that used by others.²¹

The Application of siRNA or Neutralizing Antibody

TSLP-specific siRNA reagents (SMARTpool: a mixture of 4 siRNAs; ON-TARGETplus) and negative control nonspecific siRNAs were designed by and purchased from GE Dharmacon Company (Lafayette, CO, USA). Mice were subconjunctivally injected twice with siRNAs (10 μ M, 5 μ L) over 2 days at -24 and -4 hours prior to *P. aeruginosa* infection (time 0). To apply neutralizing antibody, mice were subconjunctivally injected with rabbit-anti-TSLP (#4023, Prosci, Poway, CA, USA) and goat-anti-TSLPR (#AF546, 200 ng/5 μ L; R&D Systems, Minneapolis, MN, USA) 4 hours before the inoculation with *P. aeruginosa* on the corneas.

Clinical Examination, Bacteria Load Determination, MPO Measurement, and ELISA

Clinical examination was performed with corneal photographs and clinical scoring as described previously,²² five mice in each group. We used our previously modified methods that allowed all three assays (bacteria load, MPO determination, and cytokine ELISA measurement) to be performed with a single mouse cornea. Briefly, the corneas, five in each group, were excised, minced, and homogenized in 100 µL PBS with a micro tissue grinder (Dounce; Wheaton, Milville, NJ, USA). The homogenates were divided into two. The first part was subjected to plate bacterium counting. Aliquots (50 µL) of serial dilutions were plated onto pseudomonas isolation agar plates in triplicates, and colonies were counted the next day. The results were expressed as the mean number of CFU/ cornea \pm standard error. The second part of the homogenates was mixed with 5 μL of 1% SDS and 10% Triton X-100. For MPO assay, 30 µL homogenates was mixed with 270 µL of hexadecyltrimethylammonium bromide (HTAB) buffer (0.5% HTAB in 50 mM phosphate buffer, pH 6.0). The samples were then subjected to three freeze-thaw cycles, followed by centrifugation at 16,000g for 20 minutes. A total of 20 µL of the supernatant was mixed with 180 µL of 50 mM phosphate buffer (pH 6.0) containing 16.7 mg/mL 0,0-dianisidine hydrochloride and 0.0005% hydrogen peroxide at a 1:30 ratio in a 96-well plate. The change in absorbance at 460 nm was monitored continuously for 5 minutes in a microplate reader (Synergy2; BioTek). The results were expressed in units of MPO activity/cornea. For ELISA, 20 µL cell lysates was used. The ELISA was performed according to the manufacturer's instructions (#MTLP00; R&D Systems) with minimal three samples for each condition.

Semiquantitative and Quantitative PCR

Total RNA was extracted with an RNA extraction kit (RNeasy Mini Kit; Qiagen, Valencia, CA, USA) following the manufacturer's instructions. RNA was reversed-transcribed to cDNA with a first-strand synthesis system (DNA Thermal Cycler 480;



FIGURE 1. TSLP expression and secretion in primary human corneal epithelial cells in response to *P. aeruginosa* challenge. HCECs (p4) grown overnight in KBM were treated with heat-killed *P. aeruginosa* (MOI = 50) for different times. Cell lysates as well as culture media were collected and subjected to PCR and Western blotting analyses. (A) TSLP expression in cultured HCECs challenged with heat-killed *P. aeruginosa* for 1, 2, 4, and 8 hours was analyzed by real-time PCR. The results are representative of two independent experiments, each with three corneas. **P* < 0.05. ***P* < 0.01 (1-way ANOVA, with Bonferroni Correction). (B) The steady-state expression of TSLP in HCEC cell lysates at indicated times were analyzed by Western blott. (C) TSLP secretion was analyzed by Western blotting of the concentrated culture media of HCECs challenged with heat-killed *P. aeruginosa*. The data shown are representative of two independent experiments.

Derkin Elmer Letus, Norwalk, CT, USA). For semiquantitative PCR, cDNA was amplified with TaqMan technology (Promega, Madison, WI, USA). PCR products were subjected to electrophoresis on 2% agarose gels containing ethidium bromide. For quantitative PCR, cDNA from three corneas was amplified using a Real-Time PCR system (StepOnePlus; Applied Biosystems, University Park, IL, USA) with a PCR Master Mix (SYBR Green; Applied Biosystems). Data were analyzed by using $\Delta\Delta$ CT method with β -actin or GAPDH as the internal control.

Western Blot

Mouse corneal samples were lysed with RIPA buffer. The lysates were centrifuged to obtain supernatant. Protein concentration was determined by BCA assay with a Protein Assay Kit (Micro BCA; Pierce Biotechnology, Rockford, IL, USA). The protein samples were separated by SDS-PAGE and electrically transferred onto nitrocellulose membranes (Bio-Rad Laboratories; Hercules, CA, USA). The membranes were blocked with 3% BSA and incubated with primary antibodies overnight at 4°C. After several washes, the membranes were incubated with horseradish peroxidase-conjugated secondary antibodies. Signals were visualized using a chemiluminescent substrate (SuperSignal West Pico; Thermo Fisher Scientific, Pittsburgh, PA, USA). β -actin was used as the loading control. Antibodies: rabbit-anti-TSLP and goat-anti-TSLPR. For each condition, two samples were shown to indicate similar band intensity.

Whole-Mount Confocal Microscopy

Mice were euthanized, and the entire cornea plus the limbus was excised under the operating microscope. Excised corneas



FIGURE 2. TSLP and TSLPR expression in *P. aeruginosa*-infected mouse corneal epithelial cells. The centers of B6 mouse corneas were gently scratched with 26-gauge needles (1 mm long; 3 lines) and incubated with 1.0×10^4 CFU of *P. aeruginosa*. At the indicated times, mouse CECs were collected and subjected to (**A**) RT-PCR, (**B**, **C**) Real-time PCR for TSLP and TSLPR expression in naïve and *P. aeruginosa* infected mouse CECs, respectively. (**D**) TSLP expression in mouse CECs was assessed using ELISA. *Error bars* show standard deviations. The results are representative of two independent experiments, each with 3 corneas. *P < 0.05. **P < 0.01 (1-way ANOVA, with Bonferroni correction).

were fixed in 4% paraformaldehyde and stored at 4°C until further processing. Before staining, radial incisions were made to produce six pie-shaped wedges. Corneas were washed in PBS, incubated in 20 mmol/L pre-warmed EDTA for 30 minutes at 37°C, and incubated with a 0.2% solution of Triton X-100 in PBS plus 1% bovine serum albumin (BSA) with Fc block for 20 minutes at room temperature. After blocking, the corneas were incubated overnight at 4°C with 100 L of hamster CD11c antibody (BD Pharmingen, San Diego, CA, USA) and goat TSLPR antibody (AF546; R&D Systems) diluted in PBS with 1% BSA. The tissues were then washed five times in PBS. Corneas were then incubated with 100 L AlexaFluor 594-conjugated anti-goat antibody and FITC-conjugated anti-Armenian hamster antibody diluted in PBS with 1% BSA for 1 hour at room temperature. This was followed by five washes in PBS. Stained tissue whole mounts were placed in mounting medium (Vectashield; Vector Laboratories, Burlingame, CA, USA) onto glass slides and coverslipped. Corneal whole mounts were examined using confocal microscopy (TCSSP2; Leica, Wetzlar, Germany) for immunopositive cells in the cornea.

Depletion of DCs

CD11c-DTR and WT B6 mice were injected with 100 ng of DT in 100 μ L of PBS intraperitoneally (IP). Twenty-four hours after DT injection, the corneas were incubated with 1.0×10^4 CFU of *P. aeruginosa*; the progress of infection was monitored for up to 3 days postinjection (dpi). This procedure was shown to deplete DCs systematically of CD11c-DTR but not WT B6 mice.²³

Statistical Analysis

Data were presented as mean values \pm SD. For between-group comparisons, an unpaired, 2-tailed Student *t*-test was per-



FIGURE 3. Expression and distribution of TSLPR in residential and infiltrated DCs in naïve and *P. aeruginosa* infected mouse corneas. B6 mouse corneas (n = 5 for each condition) were inoculated with *P. aeruginosa* as described in Figure 2. Naïve (A) or *P. aeruginosa* infected, 6 hpi (B) corneas were collected for whole mount confocal microscopy and stained with TSLPR (*red*) and CD11c (*green*). Note: The CD11c-positive cells in naïve corneas can be divided into two different phenotypes: dendriform DCs (*arrowheads*) and oval ones (*arrows*). L, the limbus; C, the cornea. While dendriform DCs were TSLPR negative, Oval DCs in the limbus and naïve and infected corneas were all TSLPR positive.

formed, and for three or more conditions 1-way ANOVA with Bonferroni Correction were performed to determine statistical significance of differences in fungal counts, cytokine ELISA findings, and the MPO assay. A nonparametric Mann-Whitney U test was performed to determine the statistical significance of differences in clinical scores. Experiments with five mice for each condition were repeated at least twice to ensure reproducibility, and differences were considered statistically significant at values of P < 0.05. The results from one experiment were not compared with those of a repeated experiment because the severity of keratitis may differ among experiments. Each experiment was performed with 5 WT, untreated B6 mice as the control.

RESULTS

TSLP Expression in Primary Human CECs in Response to *P. aeruginosa* Challenge

We first investigated whether HCECs express TSLP in response to *P. aeruginosa* challenge in vitro. Primary HCECs (passage 4) were challenged with heat-killed *P. aeruginosa*. TSLP at the mRNA and protein levels were assessed by qPCR and Western blotting (Fig. 1). TSLP expression at the mRNA levels was significantly increased at both 1 and 2 hours postchallenge (Fig. 1A), followed by a rapid decline to a level slightly, but significantly higher than that in naïve cells (set as a value of 1) at 4 and 8 hours post infection (hpi; Fig. 1A). At the protein level, while TSLP was detected in the control (0 hours) and 3hour time point, the band intensity increased drastically at 8 hours post *P. aeruginosa* challenge in cultured HCEC lysates (Fig. 1B). Interestingly, TSLP protein can be detected in the culture media of primary HCECs at 3 hours post *P. aeruginosa* challenge, suggesting that TSLP induced at this time point was primarily secreted (Fig. 1B).

TSLP Expression in Mouse Corneas in Response to *P. aeruginosa* Infection

The induction of TSLP expression has been linked to allergic conjunctivitis.^{13,14} To determine whether microbial infection also causes TSLP upregulation in the cornea, we inoculated B6 mouse corneas with 1.0×10^4 CFU of *P. aeruginosa* (strain ATCC 19660) and assessed the expression of TSLP and TSLPR in CECs at different times after bacterial inoculation. RT-PCR revealed that no TSLP mRNA can be detected in naïve and 2 hpi corneal CECs (Fig. 2A) while TSLPR was detected in the naïve condition, with a slight increase in P. aeruginosachallenged CECs at 2 hpi. At 6 hpi, strong bands were seen for TSLP while TSLPR levels remained relatively high. The expression patterns of TSLP and TSLPR revealed by RT-PCR were quantitated by qPCR (Figs. 2B, 2C). Infection induced more than a 20-fold increase in TSLP transcripts at 6 hpi (Fig. 2B) whereas the expression of TSLPR increased gradually during 2 to 6 hpi (Fig. 2B). ELISA analysis revealed a sharp increase in the levels of TSLP at 6 hpi, followed by a decrease but still significantly higher that the naïve corneas at 9 hpi (Fig. 2D). Hence, P. aeruginosa infection of the cornea results in an upregulation of TSLP in CECs starting after 3 hpi.



FIGURE 4. Effects of TSLPR neutralization mediated downregulation on the severity of *P. aeruginosa* keratitis. Mouse corneas were subconjunctivally injected 5 μ L control IgG or anti-TSLP at -4 hours. At 0 hours, the corneas were inoculated with 1.0 \times 10⁴ CFU of *P. aeruginosa*. (A) Micrographs and clinical scores of the control (IgG) and TSLPR neutralizing antibody treated corneas at 1, 2, and 3 dpi. Note the increased opacification in anti-TSLPR-treated corneas (*right panels*). The infected corneas were excised, homogenized, and subjected to (B) bacteria counting, (C) MPO determination (units/cornea), and (D) ELISA determination for IL1 β . The results are representative for two independent experiments. ***P* < 0.01, (*t*-test).

TSLPR is Mostly Colocalized With CD11c Positive Cells

Having shown epithelial expression of TSLP, we next assessed the target cells of TSLP using whole mount confocal microscopy, with a focus on DCs in B6 mouse corneas. In a naïve cornea (Fig. 3A), cells in the limbal region were both TSLPR (red) and CD11c (green) positive whereas in the peripheral region of the cornea, dendriform DCs²⁴ (arrowheads) were CD11c-positive but TSLPR-negative. In this region, a few non-dendriform DCs were found to be both CD11c and TSLPR positive. In infected corneas at 6 hpi, many CD11cpositive cells migrated into the cornea with a relatively high density near the limbus (Fig. 3B). Most cells in this region had similar morphology as those in the limbus and were positive for both CD11c and TSLPR (arrows). We conclude that except for some residential DCs in the epithelium, most of the corneal DCs express TSLPR in response to *P. aeruginosa* infection.

Inhibition of TSLP/TSLPR Signaling Increases the Severity of *P. aeruginosa* Keratitis

TSLP signals through IL-7R and a unique TSLP receptor (TSLPR).²⁵ To assess the effects of TSLP signaling on P. aeruginosa keratitis, we used TSLP neutralizing antibody, TSLPR-specific siRNA, and TSLPR neutralizing antibody in the B6 mouse model. Both TSLP neutralizing antibody and TSLPR siRNA treatments augmented the severity of keratitis at 1 dpi (data not shown). Figure 4 shows the effects of TSLPR neutralizing antibody on the severity of keratitis at 1, 2, and 3 dpi. The control corneas were opacified with a clearly-visible dense area covering: 20% at 1 dpi, 70% at 2 dpi, and almost 100% at 3 dpi. In TSLPR neutralizing antibody-treated corneas, the opacification covered more than 70% of the cornea with a dense uneven opacity at 1 dpi. At 2 dpi, the entire cornea was covered with heavy opacification with a ring of relatively light opacity in between the central and peripheral region. At 3 dpi, heavy opacity covered the entire cornea with corneal



FIGURE 5. The expression of TSLP and IL-23/IL-17 in *P. aeruginosa*infected mouse corneas with or without TSLPR neutralization. Mouse corneas were subconjunctivally injected TSLPR neutralizing antibody at -4 hours. At 0 hours, the corneas were incubated with 1.0×10^4 CFU of *P. aeruginosa*. (A) Micrographs and clinical scores of the control (IgG) and TSLPR neutralizing antibody treated corneas at 1 dpi. (B) Corneas (3 in each group) were excised from the enucleated eyes at 1 dpi and subjected to Real-time PCR analysis of TSLP, IL-23, IL-17A, and IL-17C. The results are presented as fold increase with the levels of naïve corneas set as 1. The Figure is representative of two independent experiments, each with five corneas. *P < 0.05. **P < 0.01 (1-way ANOVA, with Bonferroni correction).

ulceration, melting ring, and neovascularization (reddish color), pathologies usually seen in the control corneas at day 5.^{18,26} The clinical scores revealed significantly higher disease severity in the TSLPR neutralizing antibody-treated group than that observed in the IgG injected control group (6.2 vs. 11.6 at 3 dpi; Fig. 4A). Micrographs showing pathology at 1, 2, 3 dpi from all 5 corneas, from which the clinical scores were derived, were presented as supplemental Figure S1. At 3 dpi, most corneas treated with TSLPR neutralizing antibody were about to be perforated (Supplementary Fig. S1). TSLPR neutralization resulted in a significantly higher bacterial burden $(3.37 \pm 1.84 \times 10^6 \text{ vs. } 3.31 \pm 2.28 \times 10^7 \text{ CFU}, P = 0.0198)$, neutrophil infiltration (84.57 ± 39.08 vs. 319.35 ± 28.04 units/cornea) and the expression of IL-1 β at the protein levels (811.1 vs. 3870.5 pg/µg protein) when compared to the control, IgG injected corneas at 3 dpi (Figs. 4B, 4C, 4D).

Blockade of TSLP/TSLPR Signaling Differentially Alters the Expression of IL-23 and IL-17 Cytokines in *P. aeruginosa*–Infected Corneas

Having shown that TSLP plays a protective role in *P. aeruginosa* keratitis, we next investigated whether TSLP signaling affects the expression of IL-23/IL-17 cytokines at an

earlier stage, 1 dpi since at this time point the host responses are mostly innate immunity which has grout influence on the outcome of P. aeruginosa keratitis. We used TSLPR neutralizing antibody to block TSLP signaling and observed, similar to what we had observed using TSLPR siRNA and TSLP neutralizing antibody, a significant increase in P. aeruginosa keratitis as assessed by clinical scoring, with a score of 3 in the control IgG treated and 7.5 in TSLPR neutralized corneas (Fig. 5A) at 1 dpi. As shown in Figure 5B, the infection-induced expression of TSLP was suppressed to a level lower than that in naïve corneas (from 3.1-fold increase lowered to 55.4% of the control set value as 1, a 6-fold decrease). The expression of IL-23 and its downstream cytokine IL-17A were induced by P. aeruginosa infection and further augmented by TSLPR neutralization (5.11- to 16.68- and 4.59- to 10.84-fold increase for IL-23 and IL-17A, respectively). Interestingly, TSLPR neutralization greatly downregulated IL-17c expression (from 8.48- lowered to 2.46-fold increase over the naïve corneas as the control).

Dendritic Cells Are Required for Corneal Innate Defense Against *P. aeruginosa*

Having shown that DCs are the target of TSLP, we sought to determine if DCs are required for the innate immune response for the cornea in controlling P. aeruginosa, using CD11c-DTR mice. We previously showed that subconjunctival injections of DT sufficiently depleted DCs in the cornea, resulting in significant delays in epithelial wound closure.²⁷ Our preliminary study revealed that local depletion of DCs had small effects on the outcomes of P. aeruginosa keratitis. We then used whole body depletion of DCs by 100 ng IP in 100 µi PBS. Our previous study showed this treatment depleted DCs from the spleen and cornea.²⁷ Figure 6 shows that the depletion of DCs resulted in severe keratitis, including an increase in the bacterial load $(5.84 \times 10^5 \text{ vs. } 4.40 \times 10^6 \text{ CFU} \text{ at } 3 \text{ dpi}, 13.27$ fold increase), in neutrophil infiltration as determined by MPO assay (90.0 vs. 319.35 units/cornea), and in the expression of MIP-2 (1.72 vs. 4.7 ng/cornea). The DC-depleted cornea shown at 3 dpi was about to be perforated (Fig. 6), a process normally requiring a minimum of 5 days in WT B6 mice.

The corneas with or without DC depletion were also collected at 1 dpi and subjected to qPCR analysis (Fig. 7). Unlike TSLPR neutralization which decreased TSLP and increased IL-23 expression, DC depletion had no detectable effects on infection-induced TSLP and IL-23 expression. Interestingly, the infection-induced expression of both IL-17A and -17C was significantly dampened by DC depletion.

DISCUSSION

Using the mouse P. aeruginosa-keratitis model and primary HCECs, we tested the hypothesis that TSLP plays a role in antibacterial innate immune defense. Our results demonstrate that the inoculation of P. aeruginosa stimulates both human and mouse CECs to produce and/or secrete TSLP. Downregulation of TSLP signaling with TSLPR neutralization aggravated the severity of keratitis, including great elevations of inflammation (increasing corneal opacification, clinical scores, PMN infiltration, and the expression of pro-inflammatory cytokine IL-1 β) and a moderate increase in bacterial burden, suggesting a protective role of TSLP in mediating host response to P. aeruginosa infection. Moreover, we demonstrated that most CD11c-positive cells expressed TSLPR in the infected corneas. In addition, TSLP differentially affected the expression of Th17 cytokines in B6 mouse corneas: the inhibition of TSLPR induced an up-regulation of infection-induced IL-17A but



FIGURE 6. Depletion of dendritic cells and the severe *P. aeruginosa* keratitis. CD11c-DTR and WT mice were injected with 100 ng of DT in 100 μ L of PBS IP 24 hours after DT injection, the corneas were incubated with 1.0×10^4 CFU of *P. aeruginosa*; the progress of infection was monitored for 3 days. (A) Micrographs of infected corneas at 3 dpi. Note that the DC depleted corneas (CD11c-DTR) were perforting (clinical score 12) while the control cornea (WT) remained intact with center covered by opacification. The corneas were excised and subjected to (B) a standard plate count to determine the number of bacteria in each cornea; (C) MPO determination (units/cornea); and (D) ELISA determination for MIP-2 (CXCL2). The results are representative for two independent experiments. **P < 0.01 (*t*-test).

downregulated IL-17C. Depletion of DCs resulted in severe keratitis and greatly down-regulated expression of IL-17A and C. Taken together, our results suggest that TSLP plays a protective role in *P. aeruginosa* keratitis by activating DCs and by altering the expression of IL-23/IL-17 cytokines in the cornea.

TSLP is a cytokine that is mostly epithelium-expressed, and DCs are its primary target.²⁸ At the ocular surface, the expression of TSLP was linked to allergic conjunctivitis in BALB/c mice as well as patients with various types of allergic conjunctivitis in a Th2-response related manner.^{13,14,29} In vitro, Aspergillus fumigatus stimulated the expression of TSLP in cultured human CECs in a TLR2/MyD88 dependent manner.^{15,16} In this study, we showed that challenging primary HCECs with heat-killed P. aeruginosa resulted in the upregulation of TSLP at mRNA levels within the first 2 hours, and at the protein levels at up to 8 hours. TSLP can also be detected in the culture media at 3 hpi. In vivo, P. aeruginosa infection stimulated TSLP expression between 2 and 6 hpi and peaked at the protein level at 6 hpi, followed by a decline at 9 hpi. Hence, unlike the IL-1 family of cytokines that are expressed rapidly (within first 3 hours) and continue to rise with the pathogenesis of microbial keratitis,³⁰ TSLP expression was relatively late in mouse corneas in response to P. aeruginosa infection. Importantly, the expression of TSLPR was also increased even before TSLP up-regulation in CECs, suggesting autocrine signaling of TSLP in the epithelia in response to P. aeruginosa infection.

Using antibody neutralization, we showed that downregulating TSLP-TSLPR signaling significantly augmented corneal opacification, clinical scores, bacterial burden, PMN infiltration, and the expression of IL-1 β at 3 dpi. The bacterial plate counts for the controls of *P. aeruginosa* toxic strain at 3 dpi are within the ranges reported in the literature^{31,32} as well as our previous studies,^{18,26} even though the number of bacteria used to inoculate the mouse corneas were 100-fold less than that used by others.^{21,31} The differences in the detected bacterial burden between the control and TSLPR-neutralized corneas at 3 dpi (~10 fold with P = 0.0198) is comparable to our previous study of the corneas treated with recombinant IL-24.²⁶ Together, these parameters-opacification shown in micrograph, clinical scoring, bacterial plate counting, MPO measurement, and proinflammatory cytokine detection-revealed a significant increase in the severity of keratitis in B6 mice. In the literature, TSLP has been shown to enhance neutrophil killing of methicillin-resistant Staphylococcus aureus without affecting TH2 responses during an in vivo skin infection.³³ On the other hand, overproduction of TSLP by intestinal epithelial cells negatively regulated IL-22 production by group 3 innate lymphoid cells and impaired innate immunity to Citrobacter rodentium infection.³⁴ Our study indicated a protective role of TSLP-TSLPR signaling in corneal innate immune defense against P. aeruginosa infection in B6 mice similar to that observed in the skin.³³ Disruption of TSLPR signaling resulted in a marked reduction of TSLP mRNA levels in infected corneas, suggesting a positive feedback loop of regulation of the epithelial cytokine. A positive feedback loop that drives TSLP transcription in a STAT-dependent manner was reported recently in intestinal and skin epithelial cells.35 TSLP has also been known to both induce Th2 differentiation and to be induced by activated Th2 cells, resulting in a positive feedback loop to enhance allergic inflammatory conditions.³⁶ Hence, the positive feedback loop should amplify the protective effects of TSLP in the corneas in response to P. aeruginosa infection.

We assessed the effects of TSLPR signaling blockade on the expression of cytokines with a focus on IL-23/IL17 pathways in the B6 mouse corneas. This is because TSLP targets DCs and DCs are known to be a major source of IL-23 under many homeostatic and pathologic conditions.³⁷ Moreover, TSLP was shown to induce the production of copious amounts of IL23 in primary human DCs.³⁸ Importantly, the IL-23/IL-17 immune axis has been recognized as a critical element of innate



FIGURE 7. The expression of TSLP and IL-23/IL-17 in DC depleted, *P. aeruginosa*-infected B6 mouse corneas. CD11c-DTR and WT mice were injected with 100 ng of DT in 100 µL of PBS IP 24 hours before 1.0×10^4 CFU of *P. aeruginosa* inoculation of the cornea. (A) Micrographs of the WT or CD11c-DTR mice treated with DT and inoculated *P. aeruginosa* at 1 dpi. Note that the DC depleted corneas had much high average clinical score (n = 5). (B) The corneas were excised from the enucleated eyes, homogenized, and subjected to real-time PCR analysis of TSLP, IL-23, IL-17A, and IL-17C. The results are presented as fold increase with the levels of naïve corneas (NL) set as 1. The figure is representative of two independent experiments. *P < 0.05. **P < 0.01 (1-way ANOVA, with Bonferroni correction).

immune defense against different pathogens.³⁹ For example, IL-17 is known as a strong inducer of antimicrobial peptides, particularly β -defensins⁴⁰ and S100A8/A9,⁴¹ which can prevent infection at mucosal surfaces and in the skin.⁴² IL-17A is the best-characterized member of the IL-17 family to date. Here we have shown that like IL-23, the expression of IL-17A, also referred to as IL-17 in the literature, was greatly upregulated when TSLPR signaling was disrupted. This could also suggest that TSLP plays an anti-inflammatory role in P. aeruginosa keratitis. Indeed, TSLP was found to inhibit IL-12p40, a subunit of IL-23, as well as IL-17 production in the intestine in response to Heligmosomoides polygyrus infection43 and the gutdwelling parasite Trichuris muris⁴⁴ to play a protective role in the mouse intestine. Our results that TSLP plays a protective and anti-inflammatory role in P. aeruginosa keratitis are consistent with the report that topical neutralization of IL-17 (IL-17A) promoted bacterial clearance and reduced the pathology of keratitis.⁴⁵ Surprisingly, the expression of IL-17C, a novel member of the IL-17 cytokine family, was markedly suppressed by TSLP neutralizing antibody. In human gastric mucosa, Helicobacter pylori infection was associated with a significant increase in IL-17C expression in gastric mucosa, and its role is still undetermined.⁴⁶ Our study suggests

that the ability to alter the expression of IL-17 may represent a mechanism underlying the anti-infection and protective role of TSLP in the cornea, and IL-17C, known to be expressed mostly by epithelia, may play a role in TSLP-TSLPR-mediated innate protection in the cornea.

DCs are known as the main target of TSLP⁴⁷ and TSLP acts as a link between epithelial cells and DCs.^{48,49} In the current study, we showed that DCs expressed TSLPR in the cornea. Using a well-established transgenic mouse line, we transiently depleted DCs and showed that DC depletion resulted in a much more severe keratitis. This result is in sharp contrast of a report that showed temporary depletion of CD11c positive cells triggered phagocyte accumulation in the spleen and enhances their innate bacterial killing capacity in an experimental infection model of Yersinia enterocolitica, a bacterium that causes food-borne acute and chronic gastrointestinal and systemic diseases in both humans and mice. 50 The ablation of DCs was also shown to enhance resistance to intranasal challenge with *Pneumococcal pneumonia*.⁵¹ As an immune privileged site, DCs are the only immune cells residing in the corneal epithelium, which lacks $\gamma \delta$ -T cells,⁵² while macrophages are present in the posterior corneal stroma.⁵³ The infiltrated $\gamma\delta$ T cells were shown to directly stimulate epithelial secretion of CXCL1, a chemokine that promotes recruitment of neutrophils^{54,55} and promotes a IL-17 response at the ocular surface.⁵⁶ Hence, lack of DCs may lead to the deficiency of the timely infiltration of $\gamma\delta$ T cell from the limbal region, resulting in delayed PMN infiltration, and the deficiency of IL-17 responses in the cornea,⁵⁷ resulting in its rapid clinical deterioration as seen at 3 dpi of DC-depleted corneas (Fig. 7). Indeed, unlike in TSLPR-neutralized corneas, DC-depletion resulted in the suppression of both IL-17A and 17C, although levels of TSLP and IL-23 was not affected. Although DCs are thought to be the major source of IL-23, it can also be produced by epithelial cells in response to injury, allergic challenge, and infection. $^{58-60}$ This may explain why a high level of IL-23 mRNA was detected in DC-depleted corneas. Our study is the first to link DCs in an infected tissue to the expression of IL-17 isoforms. It is important to note that the majority of IL-17 is associated with cells of the innate immune system such as neutrophils, $\gamma\delta T$ cells, and innate lymphoid cells⁶¹ and that epithelia express and secrete IL-23 to regulate intestinal epithelial cell homeostasis to limit mucosal damage.⁶⁰ The function of IL-17 as a whole, and of the individual isoforms IL-17A and IL-17C in the pathogenesis of P. aeruginosa keratitis is currently under investigation in our laboratory.

In summary, in this study we assessed the expression and function of an epithelium-produced cytokine TSLP in mediating corneal innate immune response to P. aeruginosa infection. We demonstrated a supportive role TSLP through its receptor TSLPR and its target DCs in the cornea. This protective role may be related to IL-23/IL-17 pathways and may require the participation of residential and infiltrating cells. While the requirement for neutrophils, macrophages, NK cells, and DCs have been documented, the involvement of other innate immune cells, such as yoT cells and innate lymphoid cells, in mediating the expression of IL-17 and in the innate immune defense against microbial keratitis are expected but have yet to be proven. The results shown in this study provide a link for future study to define the role IL-23/IL-17 and the involvement of innate immune cells in the pathogenesis of bacterial keratitis.

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