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CCR6+ $\gamma\delta$ T cells home to skin wounds and restore normal wound healing in CCR6-deficient mice

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DATA AVAILABILITY

Datasets related to this article can be found at doi:10.17632/n5tvt5hn5p.1, hosted at Mendeley.com.

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TO THE EDITOR

Delayed and chronic wounds, including diabetic wounds, pose an enormous burden on the healthcare system (Centers for Disease Control and Prevention 2014). Skin resident $\gamma\delta$ T cells promote wound closure by producing FGF7 and FGF10, which promote keratinocyte proliferation and wound re-epithelialization (Jameson et al. 2002). Resident $\gamma\delta$ T cells are reported to promote repair by stimulating stem cells residing in the hair follicle to multiply and traffic to the wound (Lee et al. 2017). Recently it was reported that CCR6⁺ $\gamma\delta$ T cells are recruited to a corneal epithelial abrasion via CCL20 where they promote healing (Li et al. 2011). Delayed wound healing in diabetic mice correlates with reduced numbers of recruited V γ_4 T cells and reduced IL-17A production (Liu et al. 2016). The precise mechanism by which $\gamma\delta$ T cells are recruited to the site of a wound and their importance in dermal skin healing in wild-type (WT) mice has yet to be determined.

To investigate the role CCR6 plays in dermal wound healing, we administered a 6 mm full thickness skin wound on the dorsum of WT and CCR6^{-/-} mice and measured wound area over 2 weeks. CCR6^{-/-} mice had significant larger wound areas beginning at day 6 and a 4-day delay in wound closure compared to WT mice (Fig. 1a–c). All animal experiments were approved by the Institutional Animal Care and Use Committee of the University of California, Davis and performed following the guidelines of Animal Welfare Act and Health Resource Extension Act.

To compare the numbers of $\gamma\delta$ T cells in the wounds of WT and CCR6^{-/-} mice, wounds were excised on days 2 and 5 and CD45⁺ leukocytes were enumerated by flow cytometry. On day 5, CCR6^{-/-} mice contained significantly fewer $\gamma\delta$ T cells than WT mice, and additional phenotyping revealed fewer $V\gamma_4$ T cells in CCR6^{-/-} mice (Fig. 1e). $V\gamma_4$ T cells are the predominant peripheral $\gamma\delta$ T cell subset and play a central role in infection (Huber et al. 2001) and skin graft rejection (Li et al. 2017). CCR6^{-/-} mice do not exhibit altered intraepithelial lymphocyte subpopulations (Varona et al. 2001; Mabuchi et al. 2013), so we expected that the decreased numbers of $\gamma\delta$ T cells in CCR6^{-/-} wounds was due to impaired trafficking to the wound bed. To test this, lymph nodes and blood samples from unwounded and day 2 and 5 post-wounded mice were evaluated. In both CCR6^{-/-} and WT mice, fewer $\gamma\delta$ and $V\gamma_4$ T cells were present in the lymph nodes on day 2 compared to day 0, indicating that $\gamma\delta$ T cells egress from the lymph nodes into the circulation after wounding. However, on day 5 significantly more $\gamma\delta$ T cells, including the $V\gamma_4$ subset, were observed in CCR6^{-/-} mice compared with WT mice (Fig. 1f). In blood, the frequency of circulating $\gamma\delta$ and $V\gamma_4$ T cells peaked on day 2 in CCR6^{-/-} mice before returning to levels observed in WT mice on

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day 5 (Fig. 1g). Taken together, following wounding in WT mice, $\gamma\delta$ T cells mobilized from the lymph nodes and utilized CCR6 to traffic from the circulation to the wounded skin. In contrast, following wounding in CCR6^{-/-} mice, circulating $\gamma\delta$ T cells were unable to traffic to the wound, likely due to the inability of CCR6 to interact with its chemokine ligand CCL20 (Fig. 1h).

We next sought to compare cytokine mRNA expression from day 5 wounds by RT-qPCR (Fig. 1h). CCR6^{-/-} mice had increased expression of IL-17A compared with WT mice, which is surprising given that IL-17A is produced by $\gamma\delta$ T cells in the skin of normal and diabetic wounds (Liu et al. 2016; Li et al. 2017; 2018). Other cells may be compensating for the absence of $\gamma\delta$ T cells in the wound by producing IL-17A, which can impede wound closure by inhibiting IGF-1 production by dendritic epidermal T cells (Li et al. 2018) and promote neutrophil infiltration (Takagi et al. 2017). Most remarkable was the 8-fold decrease in FGF2 expression in the wounds of CCR6^{-/-} mice compared with WT mice. FGF2 is a potent inducer of wound healing by acting on keratinocytes and fibroblasts to increase proliferation and migration at the epidermal leading edge (Kurita et al. 1992). Moreover, a subset of $\gamma\delta$ T cells exhibit the capacity to produce FGF2 (Laggner et al. 2011), raising the possibility that trafficking of CCR6⁺ $\gamma\delta$ T cells themselves might be the source of FGF2 in wounded skin. These data indicate that expression of IL-17A and FGF2 in healing skin is in part regulated by trafficking $\gamma\delta$ T cells.

We next evaluated whether normal wound healing could be restored in CCR6^{-/-} mice by adoptive transfer of WT $\gamma\delta$ T cells from WT mice. Lymph node cell suspensions from WT mice were bead-enriched for $\gamma\delta$ T cells, increasing the frequency of $\gamma\delta$ T cells from ~1% to ~85% (Fig 2a). Enriched cells were labeled with VT680 (a far-red fluorescent membrane dye) to track their trafficking by whole animal *in vivo* florescence imaging and adoptively transferred (100,000 cells/100 L) via tail vein injection into CCR6^{-/-} mice that were wounded the following day. Transferred $\gamma\delta$ T cells effectively normalized wound-healing rates in CCR6^{-/-} mice to that of the WT mice (Fig. 2b and Supplemental Fig. 1), and both conditions showed markedly faster closure than the previously wounded CCR6^{-/-} mice (Fig. 1A).

In vivo fluorescence imaging for the VT680-labeled $\gamma\delta$ T cells revealed that the exogenous $\gamma\delta$ T cells used CCR6 to traffic to wounds, which reached a maximum by 2 days and persisted through day 5 (Fig. 2c and 2d). To confirm that CCR6 was necessary for trafficking, lymph node-enriched $\gamma\delta$ T cells from WT mice were transferred into WT and CCR6^{-/-} mice and $\gamma\delta$ T cells from CCR6^{-/-} mice were transferred to WT mice. Notably, transfer of CCR6^{-/-} $\gamma\delta$ T cells to WT mice resulted in ~70% less efficient trafficking to the wound than transfer of WT $\gamma\delta$ T cells to CCR6^{-/-} mice. Furthermore, transfer of CCR6^{-/-} $\gamma\delta$ T cells to WT mice resulted in 50% less recruitment to the wound than transfer of WT $\gamma\delta$ T cells in WT mice, suggesting limited recruitment of $\gamma\delta$ T cells independent of CCR6 (Fig. 2c and 2d). Taken together, these data illustrate a critical role for CCR6⁺ $\gamma\delta$ T cells in the homeostatic process of normal wound healing.

Mobilizing CCR6⁺ $\gamma\delta$ T cells from lymph nodes into the circulation or administration of CCL20 into wounds are two potential immunological based therapies to enhance their

recruitment to impaired wounds. On the other hand, blocking CCR6 signaling is under investigation as an anti-inflammatory therapeutic in preclinical models of psoriasis and rheumatoid arthritis (Hirota et al. 2007; Mabuchi et al. 2013). We conclude that targeting CCR6/CCL20 may provide an effective strategy to maintain optimal host immune cell function and homeostasis in damaged and diseased skin.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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CONFLICTS OF INTEREST

L.S.M. has received grant support from AstraZeneca, Pfizer, Boerhinger Ingelheim, Regeneron Pharmaceuticals, Moderna Therapeutics, is a shareholder of Noveome Biotherapeutics and is on the scientific advisory board of Integrated Biotherapeutics, which are developing therapeutics and vaccines against Staphylococcus aureus and other pathogens and these associations are unrelated to the work reported in this manuscript.

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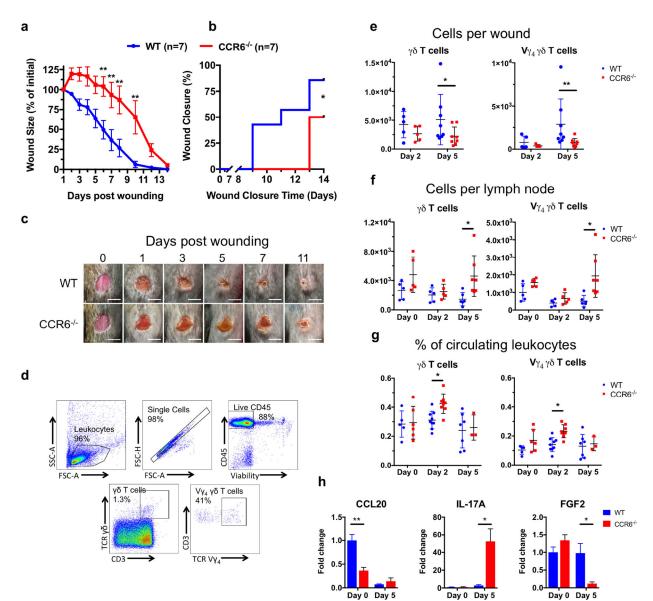


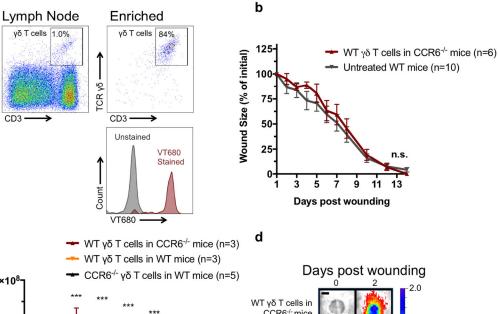
Figure 1: CCR6^{-/-} mice have hindered dermal wound healing, defective $\gamma\delta$ T cell trafficking to a wound, and dysregulated IL-17A and FGF2 expression in the wound. WT and CCR6^{-/-} mice were administered a 6mm full thickness skin wound on the dorsum. (**a-c**) Wound size was measured daily. $\gamma\delta$ T cells and V γ_4 T cells were enumerated from (**e**) wounds, (**f**) lymph nodes, and (**g**) blood by flow cytometry (**d**). Unwounded skin (day 0) or day 5 wounds were collected from animals and expression of cytokines was determined by RT-qPCR (**h**). Animals per group are presented in each graph and 4–5 animals per group for (**h**). Data is presented as Mean ± SEM.

P* 0.05 and *P* 0.01 between CCR6^{-/-} and WT. Scale bar = 5 mm.

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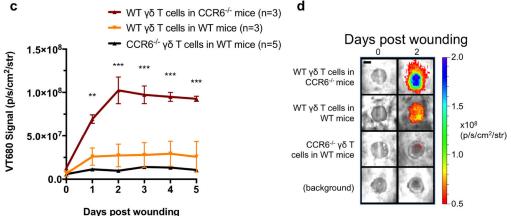


Figure 2: Adoptive transfer of $\gamma\delta$ T cells restores wound healing in CCR6^{-/-} mice. (a) $\gamma\delta$ T cells were enriched from the lymph nodes of WT and CCR6^{-/-} mice and stained with the membrane dye VT680. 100,000 $\gamma\delta$ T cells were transferred into mice via the tail vein. Mice were wounded the subsequent day, and (b) wound size was measured daily. (c and d) VT680 signal was measured for 5 days to measure $\gamma\delta$ T cell trafficking. Data is presented as Mean ± SEM. * *P****P* 0.01 and ****P* 0.001 comparing CCR6^{-/-} mice receiving WT $\gamma\delta$ T cells to other groups using a repeated measures two-way analysis of variance followed by Bonferroni's post hoc test (b and c) Scale bar = 3 mm.

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