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Supplemental information

CCL27 is a crucial regulator of immune homeostasis of the skin and mucosal tissues

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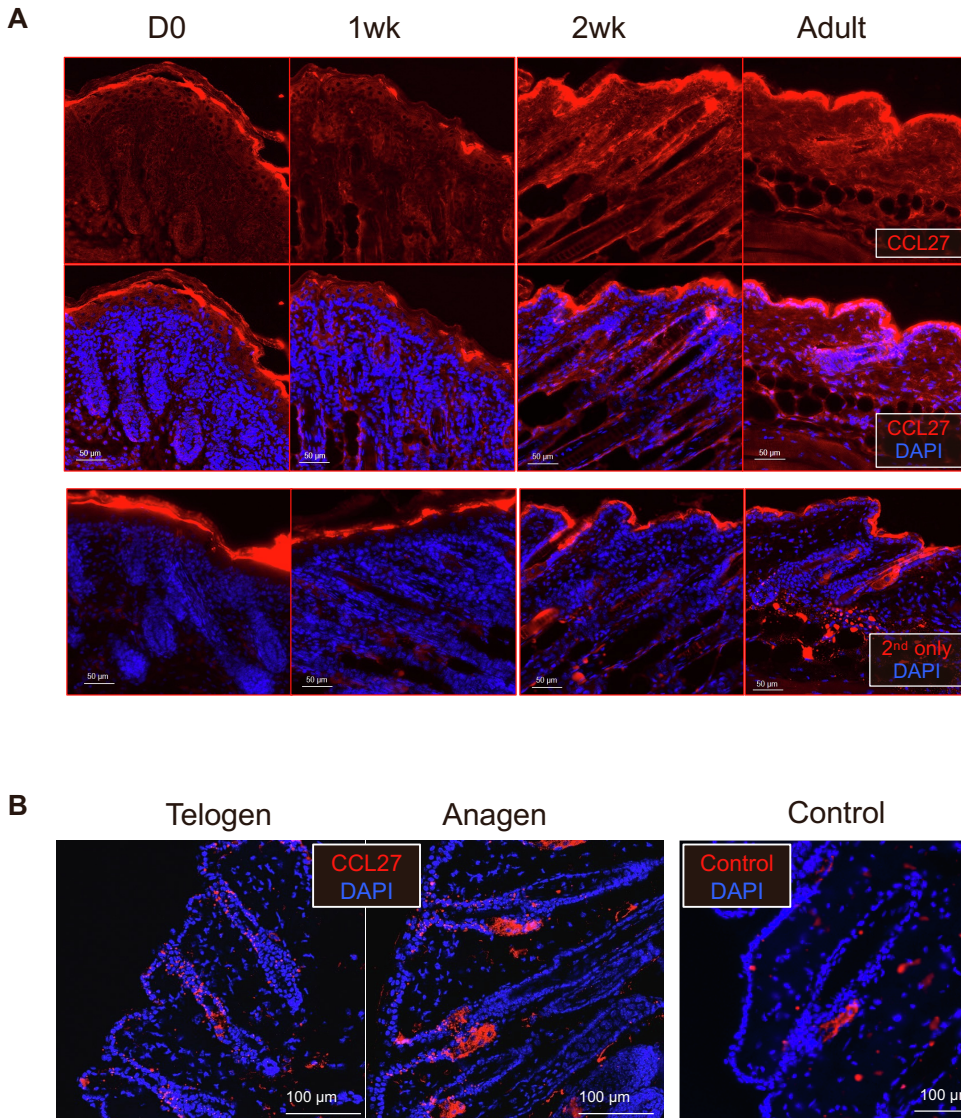


Figure S1. Preferential localization of CCR10⁺ lymphocytes near anagen phase hair follicles correlates with high CCL27 expression by follicular keratinocytes, related to Figure 1. **A)** Representative immunofluorescent skin sections (14 μ m) stained with anti-CCL27 antibody. N=7 mice for D0, 1 week, and adult ages and 6 mice for 2 weeks. **B)** Fluorescent images (7 μ m) representative of skin at resting (telogen) and growing (anagen) phase of hair follicle cycling stained by *in-situ* hybridization with an antisense CCL27 RNA probe or a non-specific RNA probe. The control probe recognizes the DapB gene (accession # EF191515) of a soil bacterial strain *Bacillus subtilis* SMY. N=3 mice per HF cycle phase. Large red patches are non-specific signals of sebaceous glands that are found in both CCL27 and control probe-stained sections.

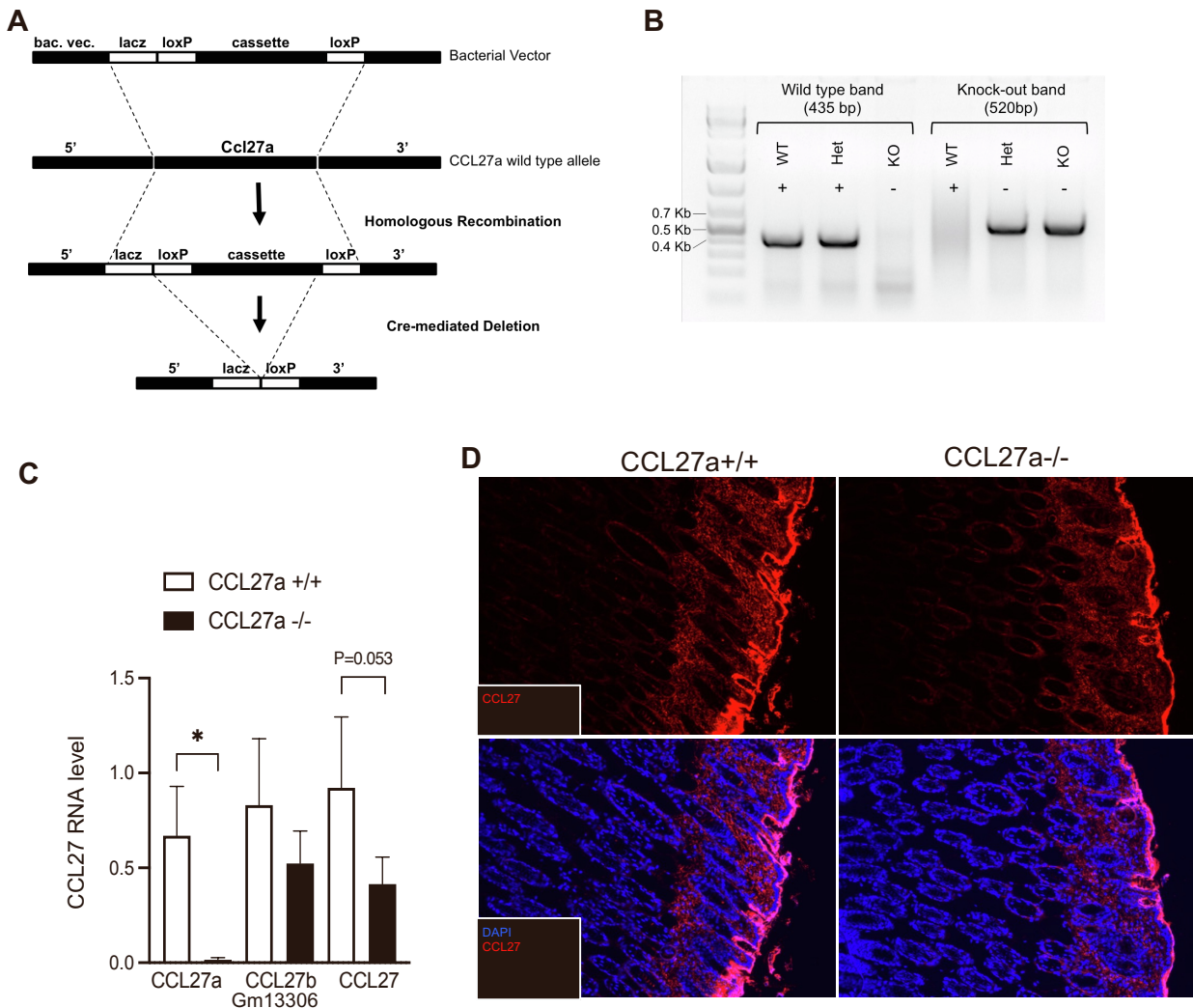


Figure S2. CCL27a-knockout has little effect on the establishment of CCR10⁺ lymphocytes in the skin, related to Figure 2. **A)** Diagram of the CCL27a-targeting strategy in the CCL27a-knockout mouse line, modified from information obtained from the KOMP Repository (https://www.mmrrc.org/catalog/sds.php?mmrrc_id=46967). **B)** Genomic PCR genotyping CCL27a^{+/+} (WT), heterozygous CCL27a^{+/-} (Het), and homozygous CCL27a^{-/-} knockout (KO) mice. Band sizes of CCL27a-WT and CCL27a-KO alleles are 435bp and 520bp respectively. **C)** Representative immunofluorescent skin sections (15µm) from CCL27a^{-/-} and CCL27a^{+/+} mice stained with anti-CCL27 antibody. **D)** Real-time RT-PCR analysis of total CCL27, CCL27a, and CCL27b/Gm13306 expression in the skin of CCL27a^{-/-} and CCL27a^{+/+} mice. Normalized to β-actin. N=6 mice for each genotype. The predicted coding sequences of CCL27b and Gm13306 are the same.

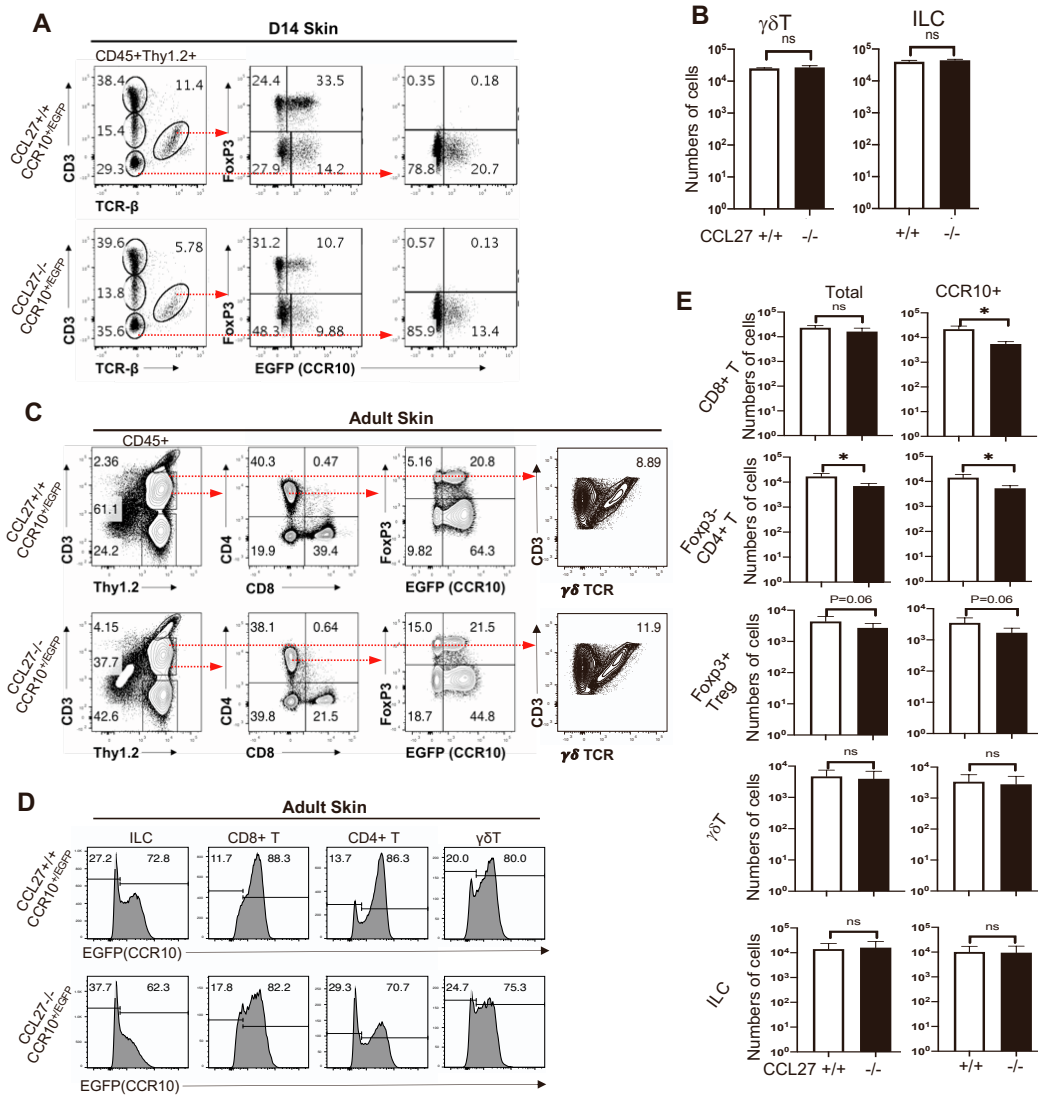


Figure S3. Dysregulated presence of CCR10⁺ lymphocytes in the skin of total CCL27-knockout mice, related to Figure 4. **A-B)** FC analysis (A) of skin lymphocytes (CD45⁺Thy1.1⁺) for numbers of different T cell subsets and ILCs (B) in two week-old CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. αβT cells are CD3^{int}TCRβ⁺. Treg cells are CD3^{int}TCRβ⁺Foxp3⁺. ILCs are CD3⁻TCRβ⁻. γδT cells are CD3^{int}TCRβ⁻ (excluding the CD3^{high} dendritic epidermal γδT cells). N=5 for CCL27^{+/+} and 4 for CCL27^{-/-} mice. **C)** FC analysis of skin CD45⁺ immune cells of adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice for different subsets of T cells and ILCs. **D)** Histograms of CCR10(EGFP) expression in the gated populations of skin T cell subsets and ILCs of adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. CD8⁺ and CD4⁺ T cells are gated on CD8⁺ and CD4⁺ CD45⁺Thy1.2⁺CD3^{int}. ILCs are CD45⁺Thy1.2⁺CD3⁻. γδT cells are CD45⁺Thy1.2⁺CD3^{int}TCRγδ⁺ (excluding CD3^{high} dendritic epidermal γδT cells). **E)** Numbers of total and CCR10⁺ CD8⁺, Foxp3⁻CD4⁺, Foxp3⁺Treg, γδ T cells and ILCs in adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. N=11 samples each for total CD8⁺ T cells, 5 each for CCR10⁺ CD8⁺ T cells, 5 each for Foxp3⁻CD4⁺ and Foxp3⁺Treg cells, and 4 each for γδT cells and ILCs.

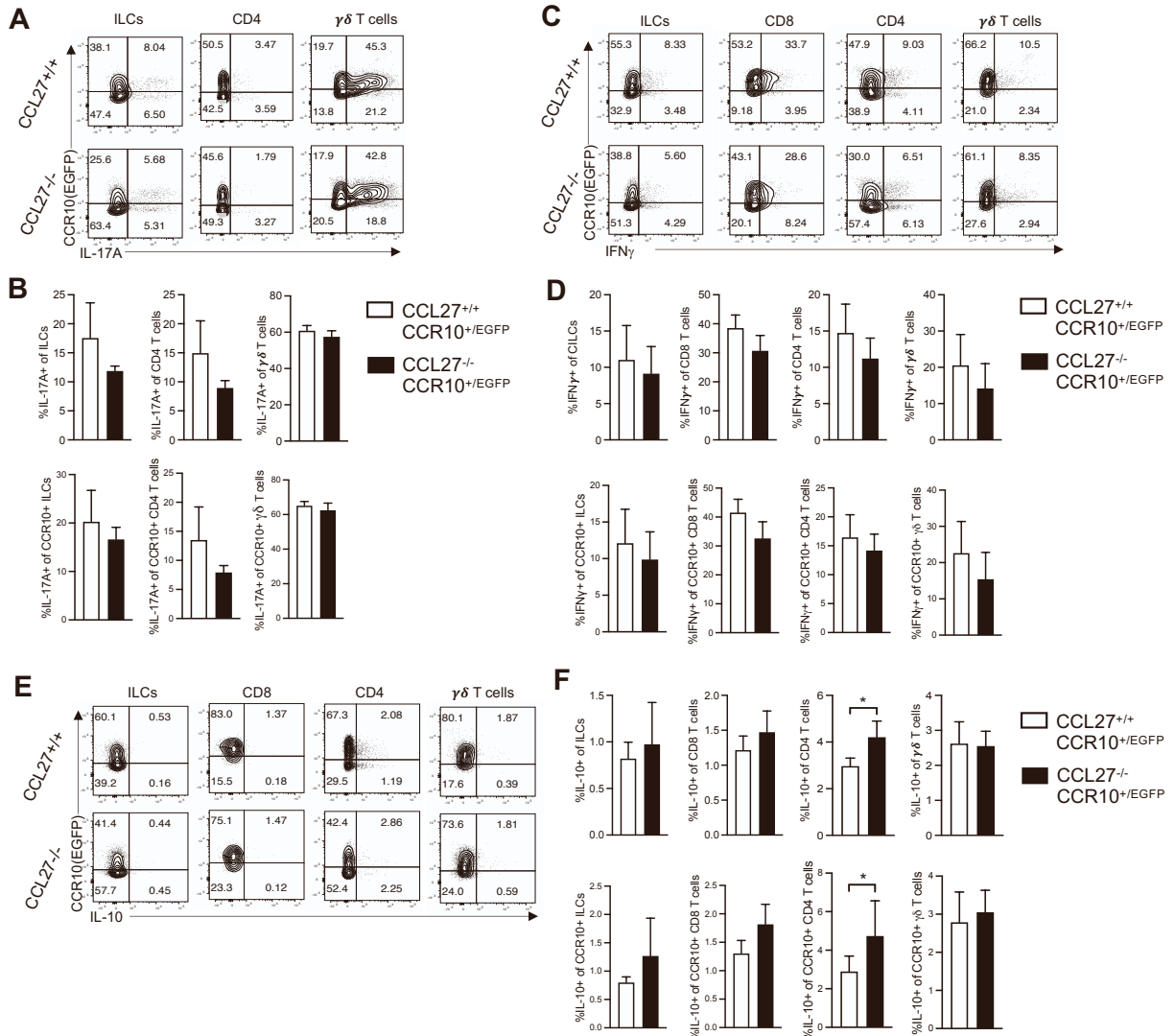


Figure S4. Minor shifts in the cytokine production capacity of skin lymphocytes in CCL27^{-/-} versus WT mice at steady state, related to Figure 4. A-B) FC analysis (A) and graphed percentages (B) of IL-17A production of stimulated skin lymphocytes from adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. N=7 mice for each genotype. C-D) FC analysis (C) and graphed percentages (D) of IFN γ production of stimulated skin lymphocytes from adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. N=6 mice each. E-F) FC analysis (E) and graphed percentages (F) of IL-10 production of stimulated skin lymphocytes from adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. T cell populations are gated on CD45⁺CD3^{int}CD8⁺, CD4⁺ or $\gamma\delta$ TCR⁺. ILCs are gated on CD45⁺Thy1.2⁺CD3⁻. N=5 mice for each genotype.

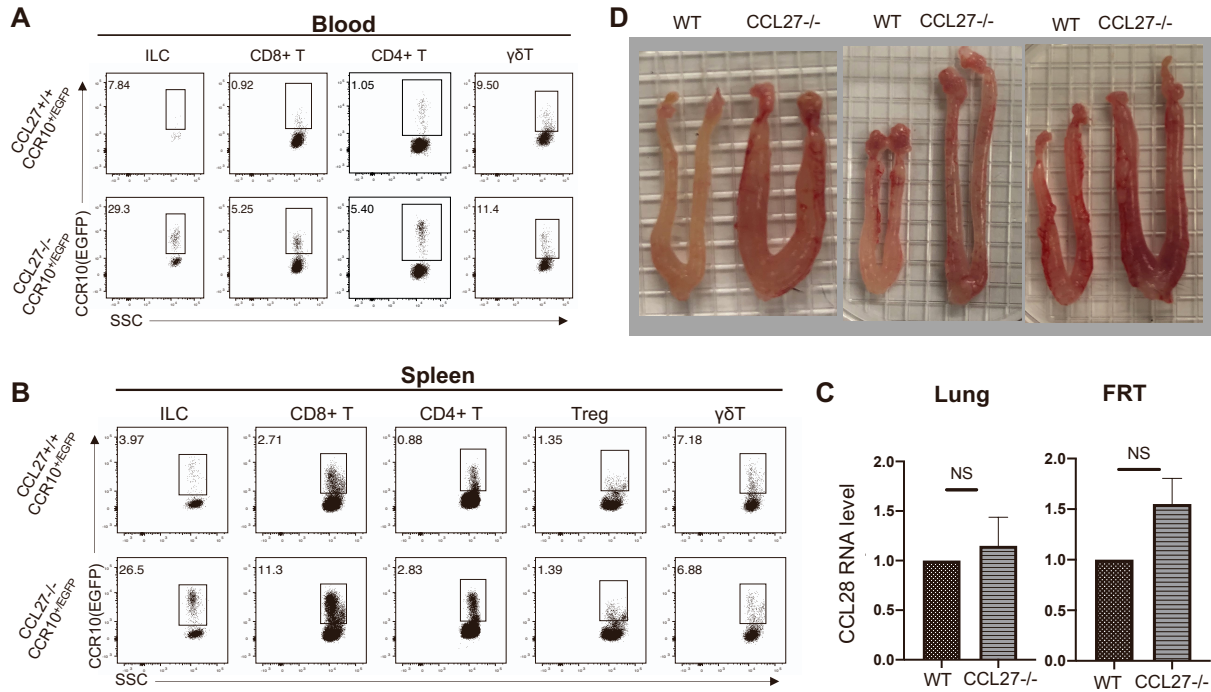


Figure S5. Increased accumulation of CCR10⁺ lymphocytes in the blood, spleens and lungs and female reproductive tracts of total CCL27-knockout mice, related to Figure 6. A-B) FC analysis gating on populations of T cell subsets and ILCs for their CCR10(EGFP) expression in the spleen (A) and blood (B) of adult CCL27^{+/+}CCR10^{+/EGFP} and CCL27^{-/-}CCR10^{+/EGFP} mice. **C)** Real-time RT-PCR analysis of relative CCL28 expression in lungs and female reproductive tracts (FRT) of CCL27^{-/-} mice in comparison to WT controls. Normalized to β-actin. N=4 each for lungs and 3 each for FRT. **D)** Images of uteri of three pairs of WT and CCL27^{-/-} female littermate mice.