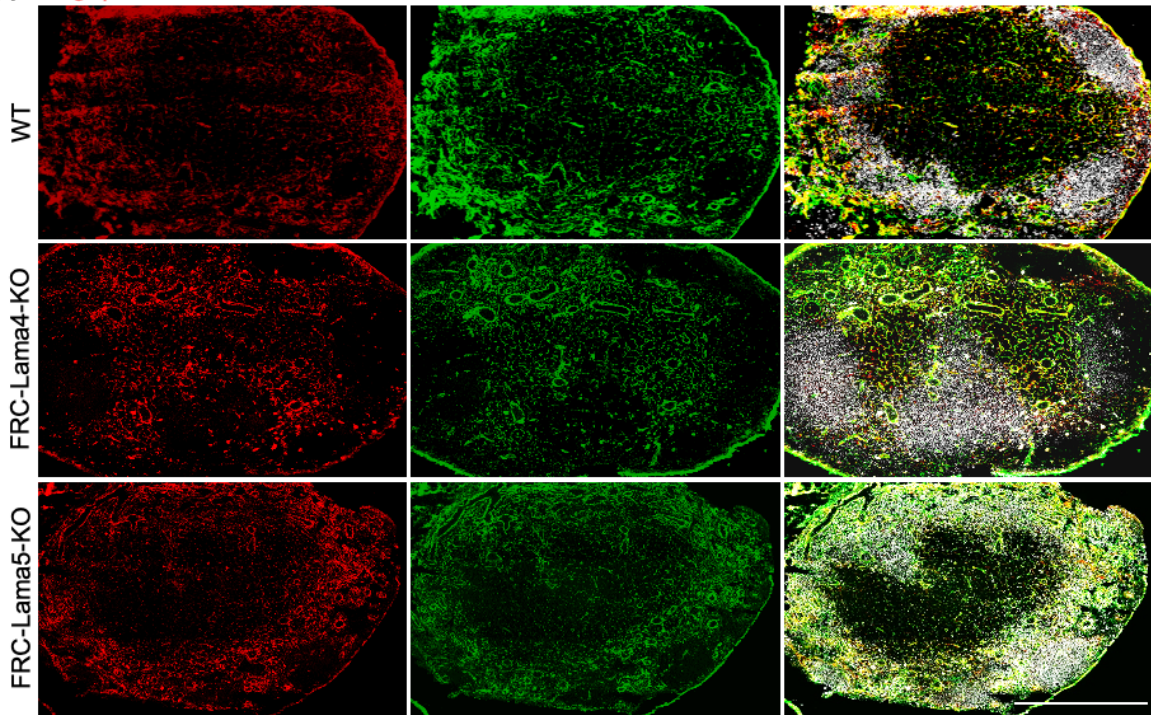
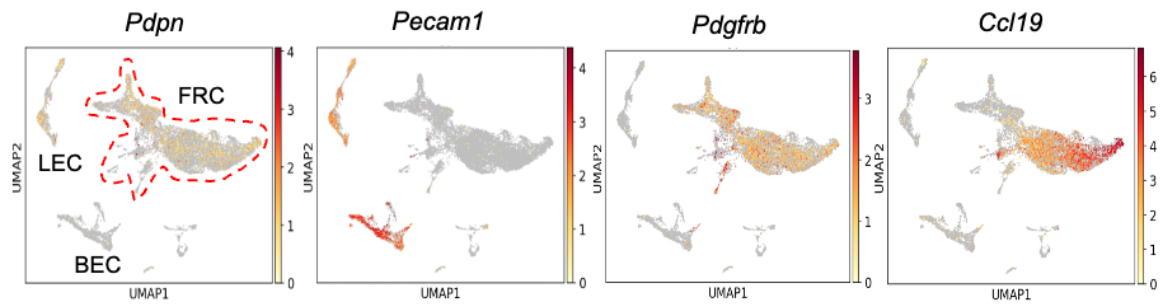


A *Pdgfrβ*/ER-TR7/B220



B



C

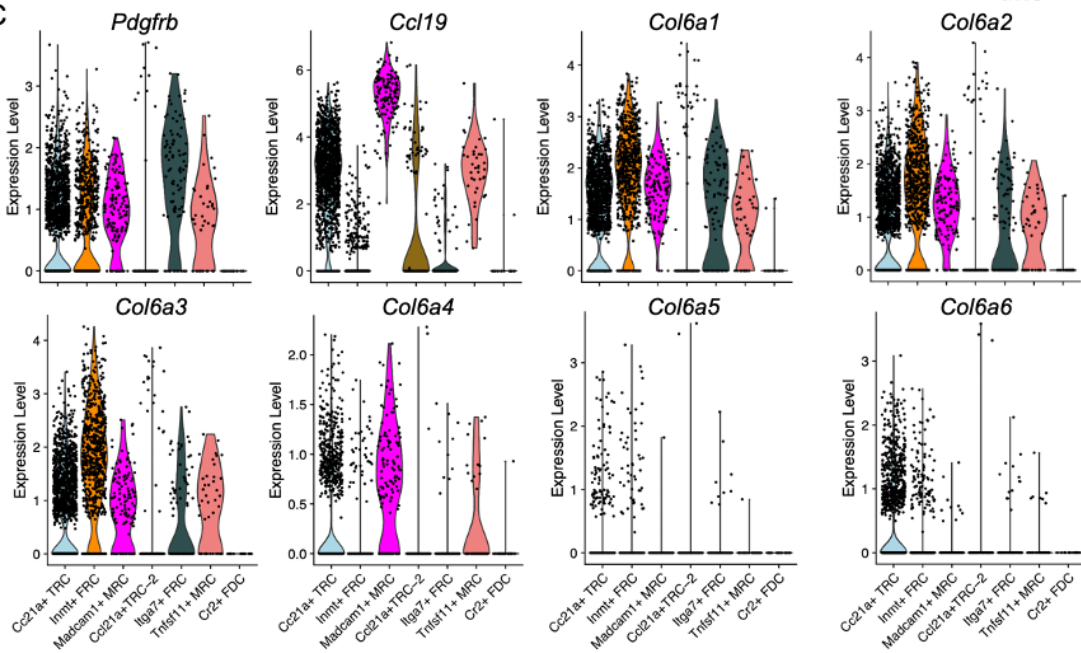


Figure S1. Platelet-derived growth factor receptor β (PDGFR β) is stably expressed by FRCs. LNs were isolated from *Pdgfrb-Cre^{-/-}* x *Laminin α 4^{fl/fl}* (WT), *Pdgfrb-Cre^{+/-}* x *Laminin α 4^{fl/fl}* (FRC-Lama4-KO), and *Pdgfrb-Cre^{+/-}* x *Laminin α 5^{fl/fl}* (FRC-Lama5-KO) mice. **A:** Whole mount scanning (20x) of 6 μ m sections stained for *Pdgfr β* , ER-TR7, B220, scale bar 500 μ m. Representative of three independent experiments. 3 mice/group, 5 LNs/mouse, 3 sections/LN, 3-5 fields/section. **B:** *Pdpn*, *Pecam1*, *Pdgfrb*, and *Ccl19* gene expression in LN stromal cells, *Ccl19* and *Pdgfrb* are exclusively expressed by *Pdpn⁺Pecam1⁻* FRCs. **C:** *Pdgfrb*, *CCL19*, *Pdpn*, and *Col6a1-Col6a6* expression in FRC subsets. FRCs were further classified into two *Ccl21⁺* T zone reticular cells (TRCs), *N-methyltransferase (Inmt)⁺* FRCs, *Madcam1⁺* marginal reticular cells (MRCs), *integrin α 7 (Itga7)⁺* FRCs, *tumor necrosis factor ligand superfamily member 11 (Tnfsf11)⁺* MRCs, and *complement receptor 2 (CR2)⁺* FDCs according to the expression of distinctive genes. Representative of three independent experiments. 3 mice/group, 5 LNs/mouse, 3 sections/LN, 3-5 fields/section.

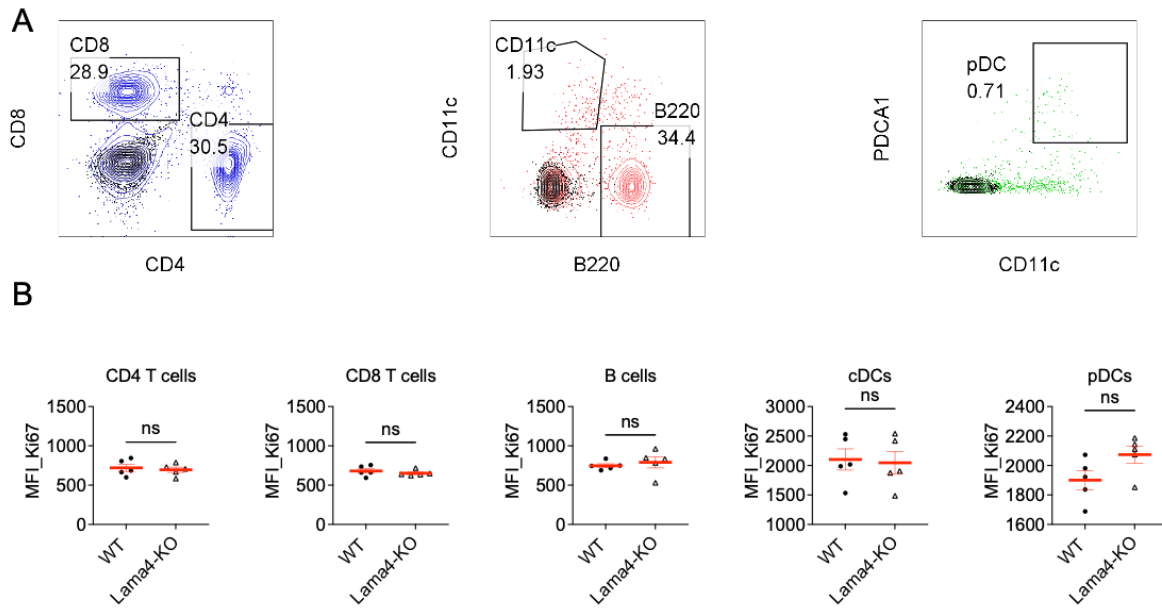


Figure S2. Proliferation of freshly isolated leukocytes. A: Gating of CD4 T cells, CD8 T cells, B cells, conventional dendritic cells (cDCs), and plasmacytoid DCs (pDCs) for flow cytometry. **B:** Ki67 intensity in these freshly isolated immune cells from WT and FRC-Lama4-KO LNs. Students' t-test for two groups comparison. Mean \pm SEM.

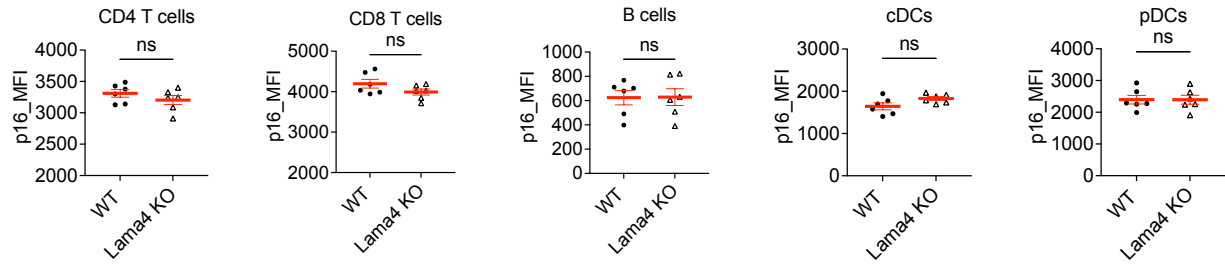


Figure S3. Senescence of freshly isolated leukocytes. Flow cytometry analysis of p16 expression in CD4 T cells, CD8 T cells, B cells, cDCs, and pDCs freshly isolated from WT, FRC-Lama4-KO, FRC-Lama5-KO LNs. Gating as in Fig. S2. Students' t-test for two groups comparison. Mean \pm SEM.

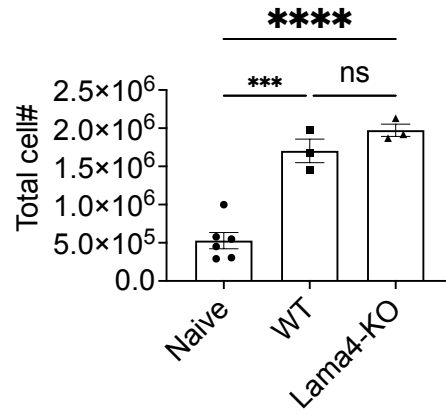


Figure S4: Mice were immunized in the flanks with IFA/OVA s.c.. Five days later, inguinal dLNs were harvested and total number of cells were quantified using flow cytometry. Representative data of two independent experiments, 3 mice/group. One-way ANOVA for multiple groups comparison. Mean \pm SEM, **** p <0.0001.

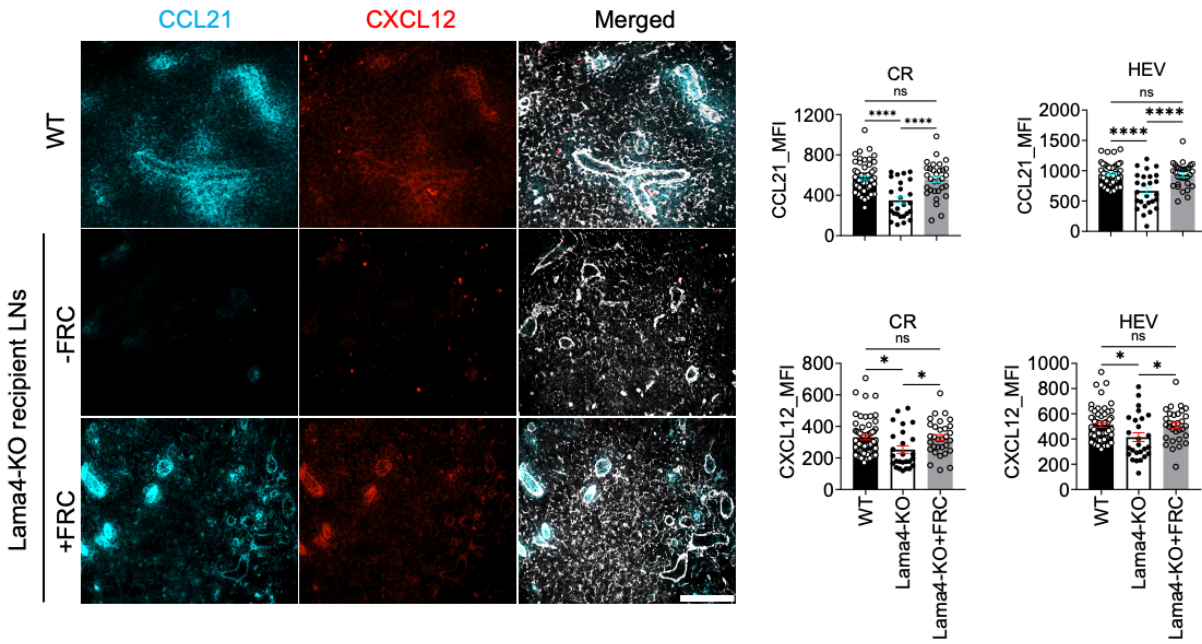


Figure S5: Transferring WT FRCs recovers chemokine distribution in FRC-lama4-KO mice. WT FRCs were injected (i.v.) into FRC-Lama4-KO mice (10^5 FRCs/dose/week x 4 weeks). LNs from WT and FRC-Lama4-KO mice with or without FRC transfer were harvested for immunofluorescent staining for CCL21 and CXCL12. Scale bar 100 μ m. Representative of three independent experiments. 3 mice/group, 5 LNs/mouse, 3 sections/LN, 3-5 fields/section. One-way ANOVA with Tukey's multiple comparisons test for multiple groups comparison. Mean \pm SEM, * $p < 0.05$, **** $p < 0.0001$.

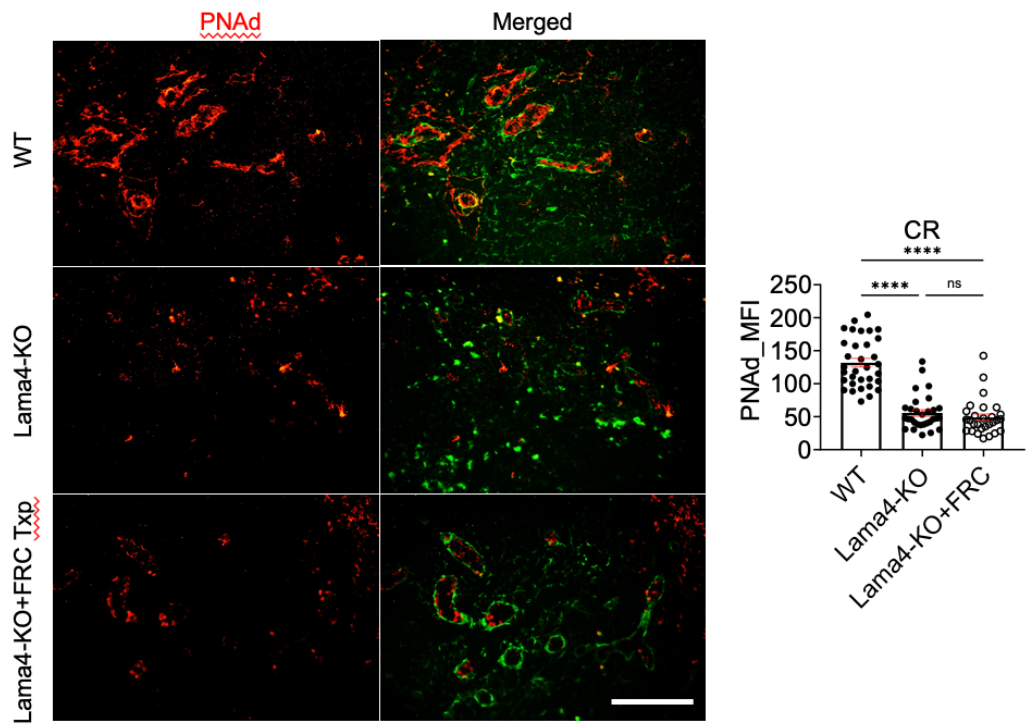


Figure S6. Transfer of WT FRCs did not rescue the defect of HEVs in FRC-Lama4-KO mice. 10^5 primary WT FRCs injected i.v. into WT and FRC-Lama4-KO mice, once a week for four weeks. One week after the 4th dose, LNs harvested. Left, cryosections stained for PNAAd and ER-TR7. Right, quantification of PNAAd intensity in cortical ridge (CR). One-way ANOVA for multiple groups comparison. Scale bar 100 μ m. Mean \pm SEM, **** $p < 0.0001$.