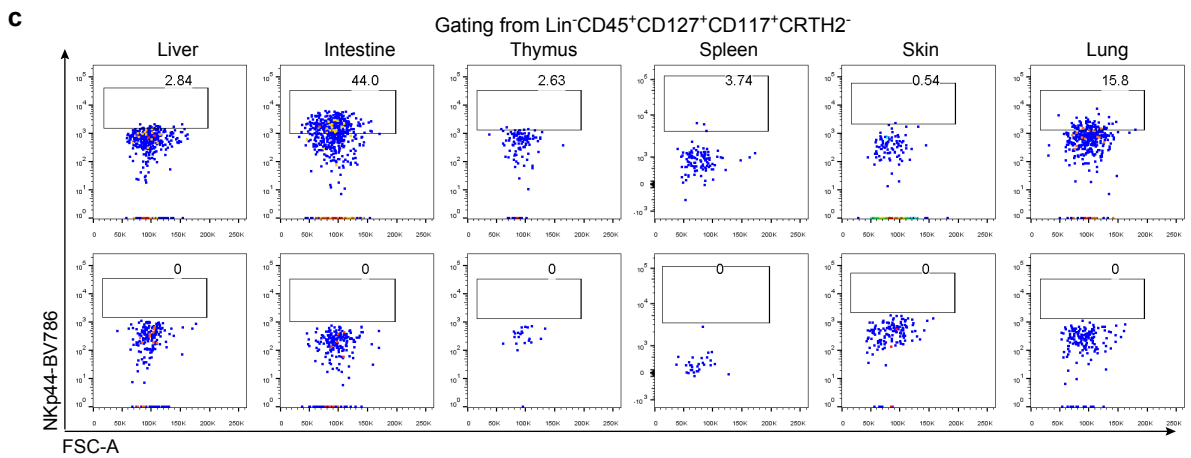
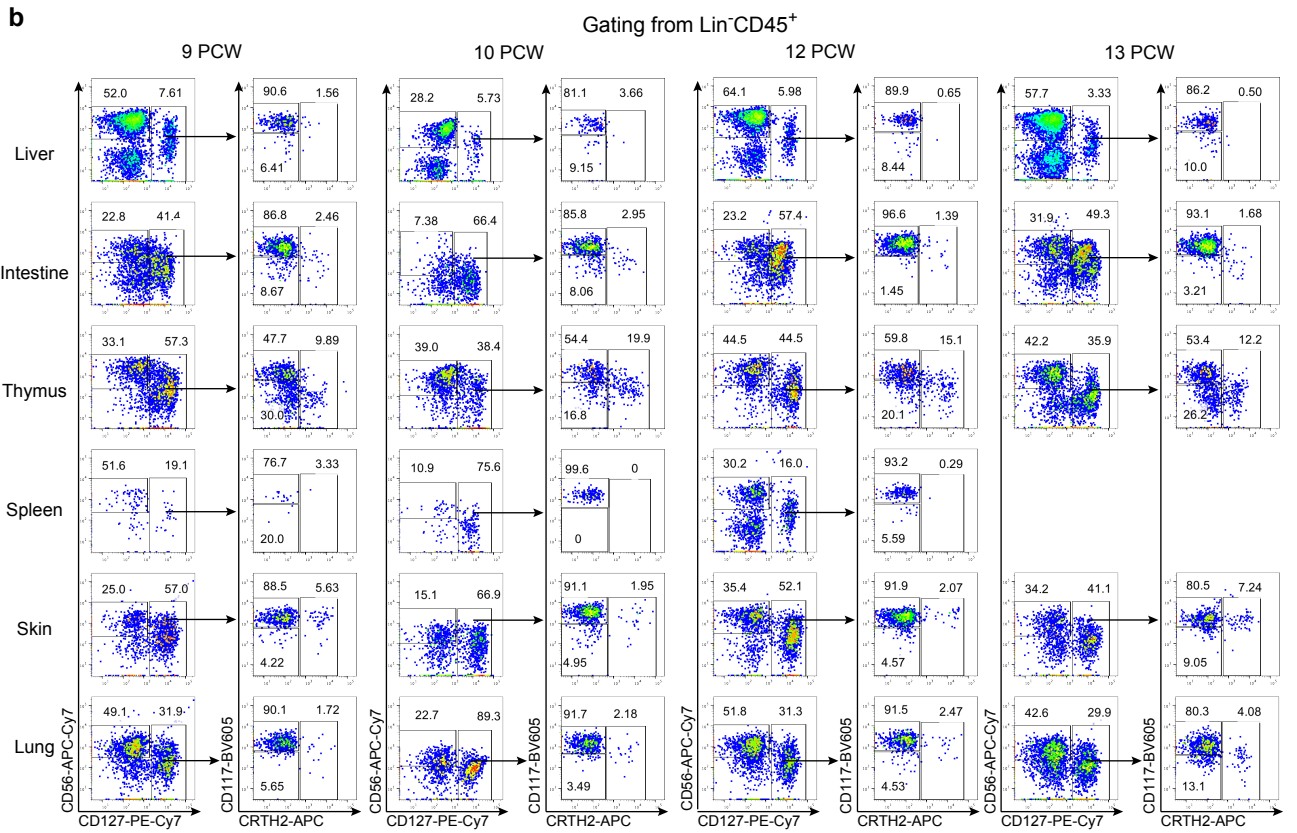
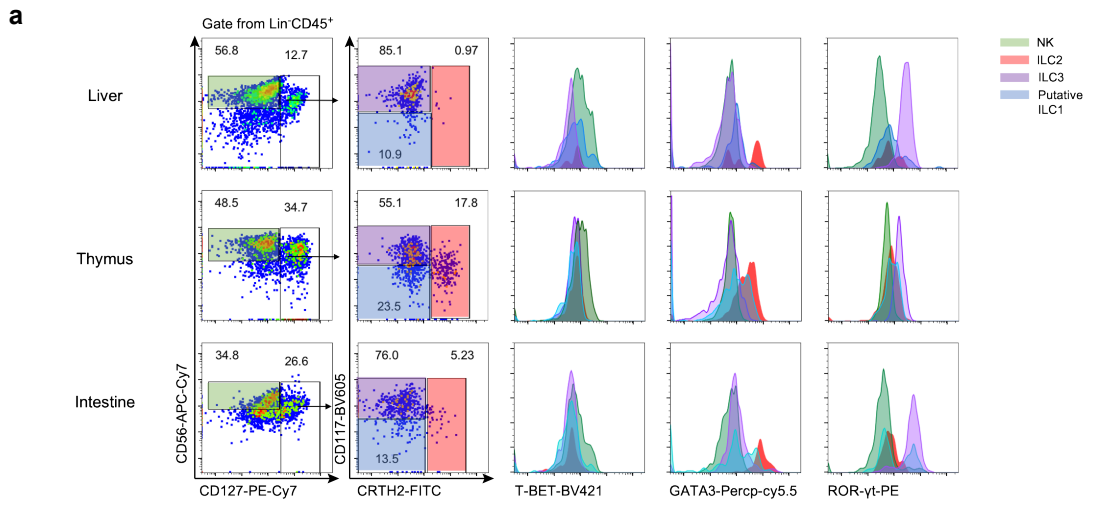


Figure S2



Supplementary Figure 2 (related to Figure 1) The tissue distribution of mature ILCs in different tissues at different developmental stages is shown by flow cytometric analysis.

(a) Expression of intracellular transcription factors (T-BET, GATA-3, and ROR γ t) in NK, ILC2, ILC3 and putative ILC1 in different tissues.

(b) Representative flow cytometry results of killer ILCs and helper ILCs subpopulations in different tissues at each stage. **(c)**

Expression of NKp44 were detected in CRTH2⁻CD117⁺ ILC3. Flow cytometric analysis of NKp44 protein level of CRTH2⁻CD117⁺ ILC3 from each tissue (upper panel) and corresponding fluorescence minus one controls (below panel).