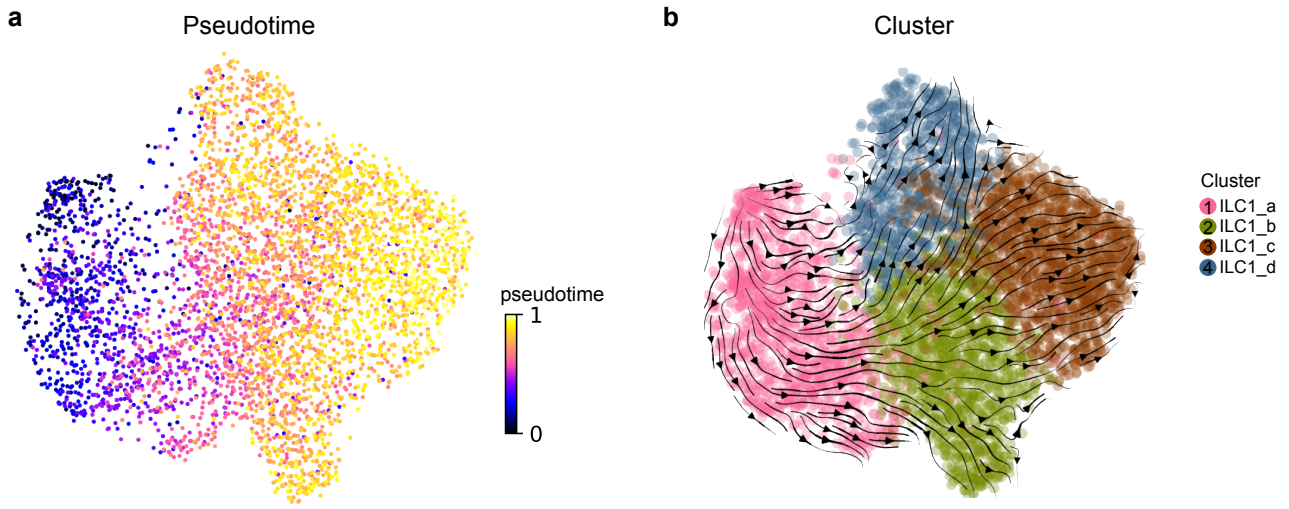


Figure S7



**Supplementary Figure 7 (related to Figure 4) Pseudotime analysis of ILC1 sub-clusters**

pseudotemporal ordering **(a)** and directed transition probabilities **(b)** were evaluated by Cellrank based on KNN graph and CytoTRACE score derived from gene expression data and projected onto umap embedding.