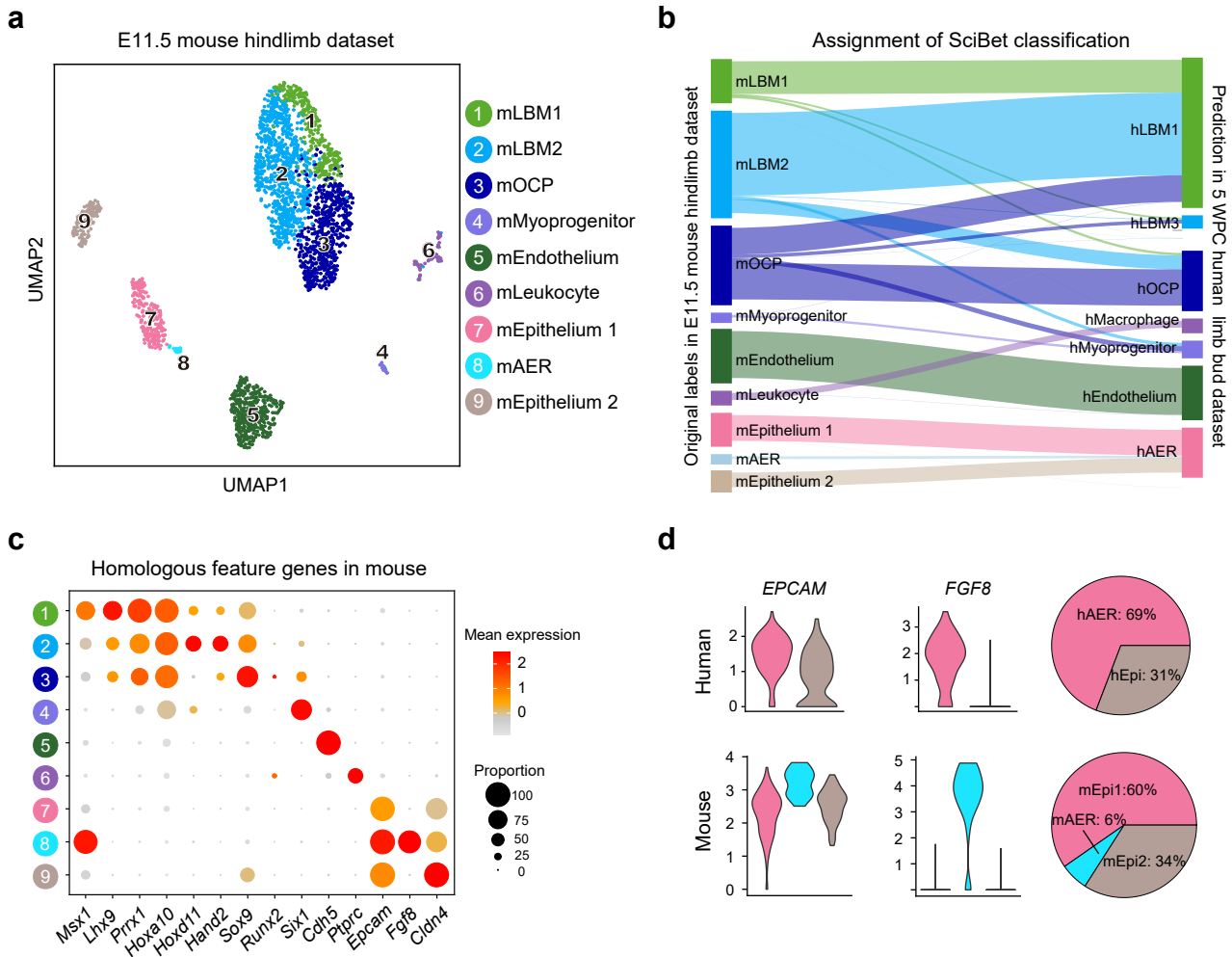


Figure S2



Supplementary Fig. 2. Characterization of E11.5 mouse hindlimb bud mesenchyme and epithelium

a, UMAP visualization of 9 cell subsets in E11.5 mouse hindlimb bud dataset. Expression matrix was re-processed and cells were clustered according to the expression of homologous feature genes in human limb bud.

b, Sankey diagram for assigning E11.5 mouse hindlimb bud subsets to 5 WPC human limb bud datasets.

c, Dot plots of mean expression of homologous feature genes in E11.5 mouse hindlimb bud subsets.

d, Violin plots (left) showing the gene expression of *EPCAM* and *FGF8* in human and mouse epithelial subsets. Pie charts (right) showing the proportions of each epithelial subset.