



Figure 3 - figure supplement 1. Unsupervised clustering of single-cell RNAseq data identifies club and basal cells. **(A).** Heatmap shows cell-cell correlation, Pearson's R (color bar) between all 74 cells in the dataset. Dendrogram (top) shows the assignment of cells to clusters (color legend). **(B).** Violin plots show the expression of basal cell marker *Krt5* (left), club cell marker *Scgb1a1* (center) and the number of genes detected per cell (right), which were used to interpret the clusters.