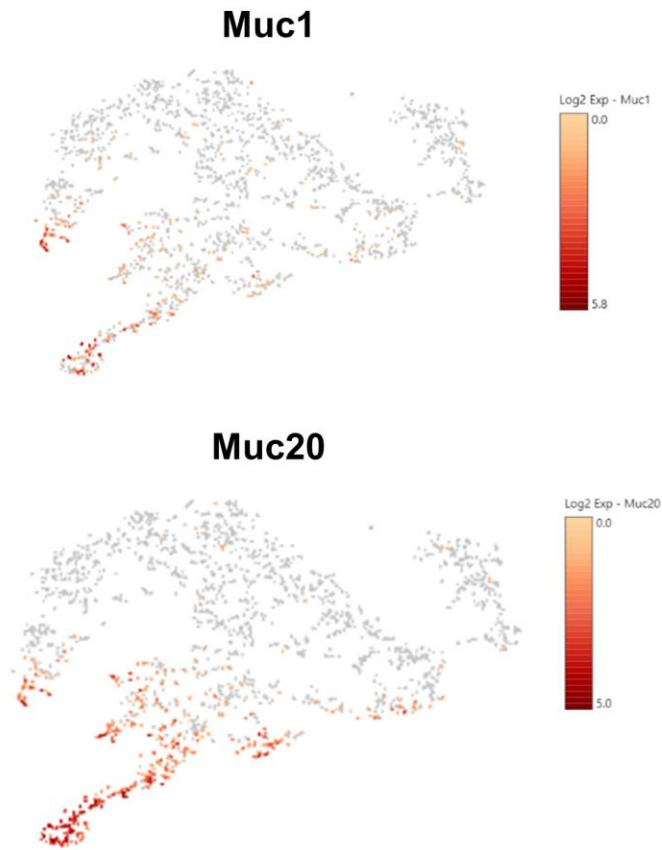


Supplementary Figure S2



Supplementary Figure S2. t-SNEs colored by the normalized log-transformed expression of conjunctival epithelial markers *Muc1* and *Muc20*. Raw sequencing data was analyzed using 10X Genomics data analysis pipeline. Image was exported via Loupe program.