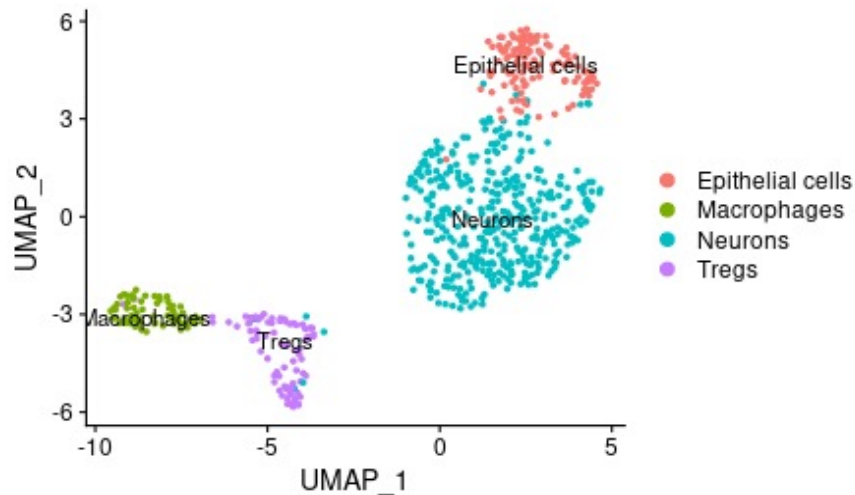
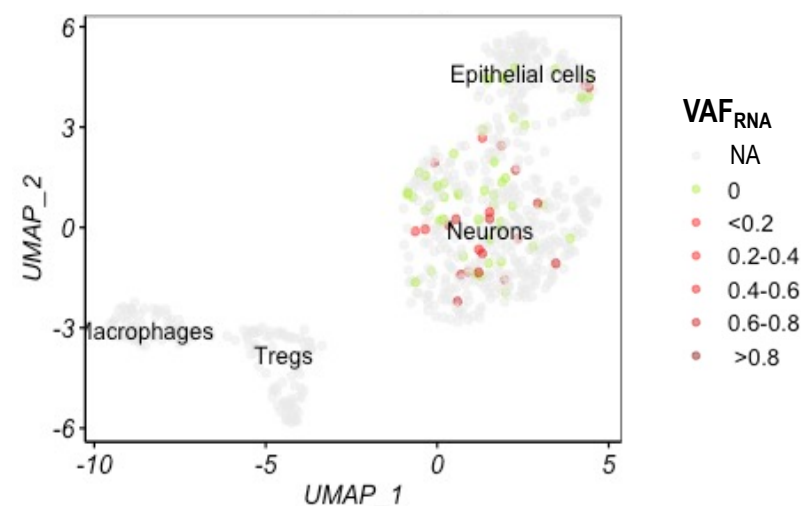


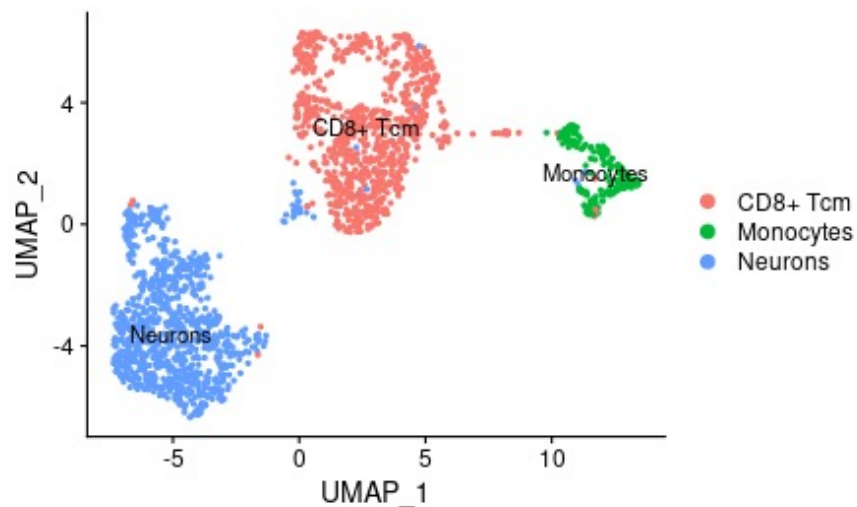
SRR10156297



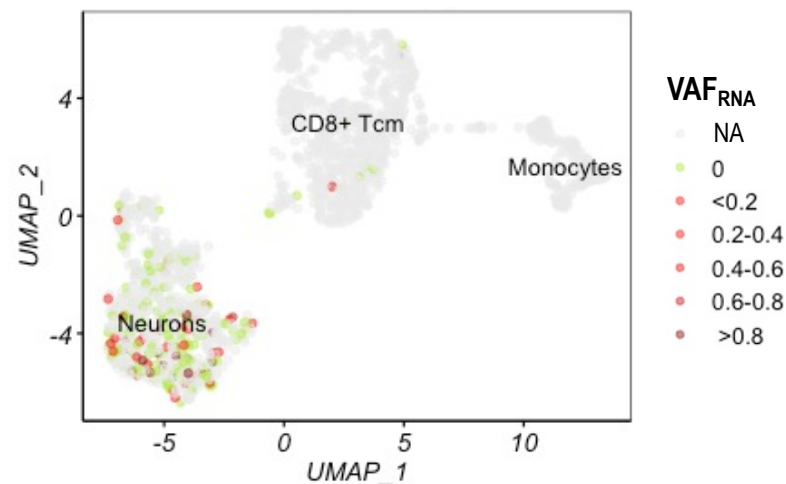
14:100846310, *MEG3*



SRR10156299



14:100846310, *MEG3*



Supplementary Figure 2. Two-dimensional UMAP clusters of samples SRR10156297 and SRR10156299 showing cells classified by type (left) and visualizing RNA-editing levels (right) in the gene *MEG3*, where the intensity of the red color corresponds to the proportion of edited reads, and the green color indicates that all the reads (minR=3) covering the position in the cell carried the reference nucleotide.