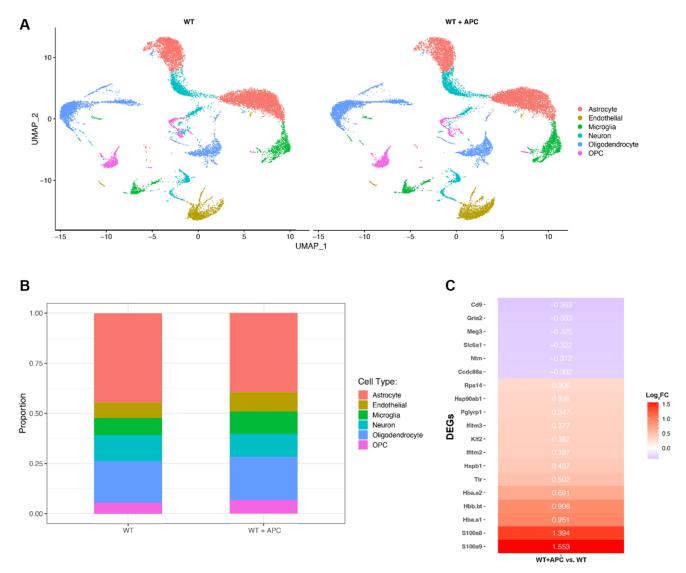
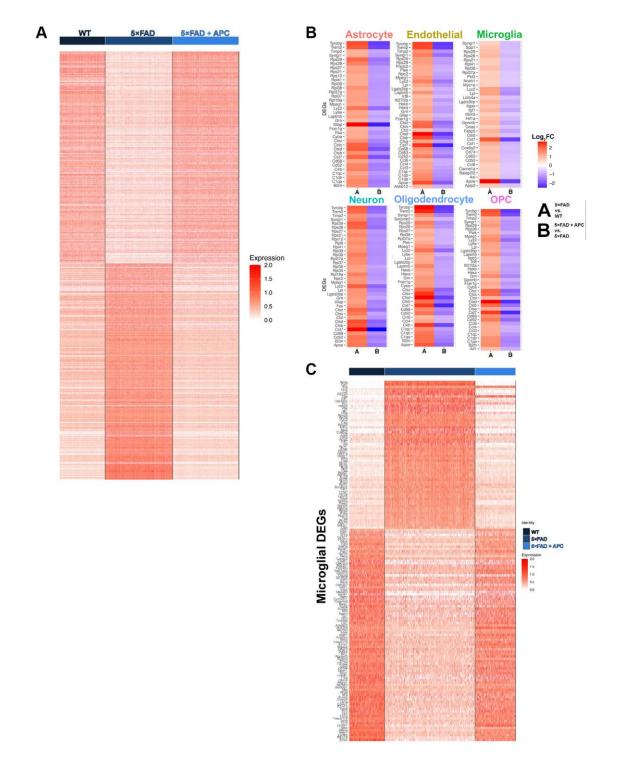
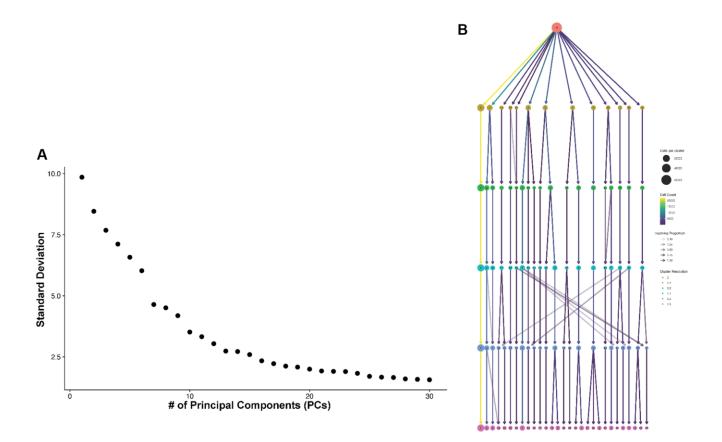
## **SUPPLEMENTARY FIGURES**



**Supplementary Figure 1.** (A) UMAP dimensional plot of control dataset: WT + APC versus WT. (B) Proportion of each cell-type observed in each sample of the control dataset. (C) Heatmap of all DEGs globally in the control dataset (*p*-value < 0.05).



**Supplementary Figure 2.** (A) Global heatmap of all DEGs in the primary dataset (~500 DEGs) (*p*-value < 0.05). (B) Heatmap of top ~30 DEGs found in every cell-type in the primary integrated dataset. (C) Heatmap of all DEGs found within the microglial subset (*p*-value < 0.05).



Supplementary Figure 3. (A) Elbow plot of variance per PC in the primary integrated dataset. (B) Cluster Tree of unsupervised clustering performed by Seurat.