

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Differential gene markers of every cluster compared to all clusters. Seurat was used to identify the differential genes expressed of every cluster compared to all clusters. Avg_logFC represents log2 fold change

File Name: Supplementary Data 2

Description: Differentially expressed genes identified from pairwise comparison of each cluster. Seurat was used to identify the differential genes expressed from a pairwise comparison of each cluster. Avg_logFC represents log2 fold change

File Name: Supplementary Data 3

Description: Seurat and Monocle cluster membership and overlap. The number of cells and frequency of overlap between the Monocle and Seurat clusters are listed