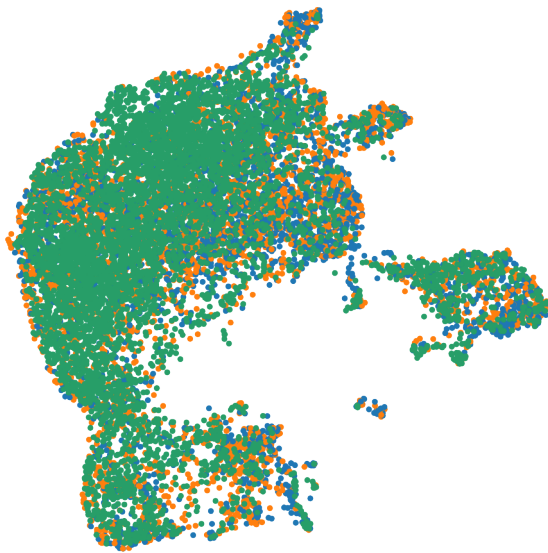
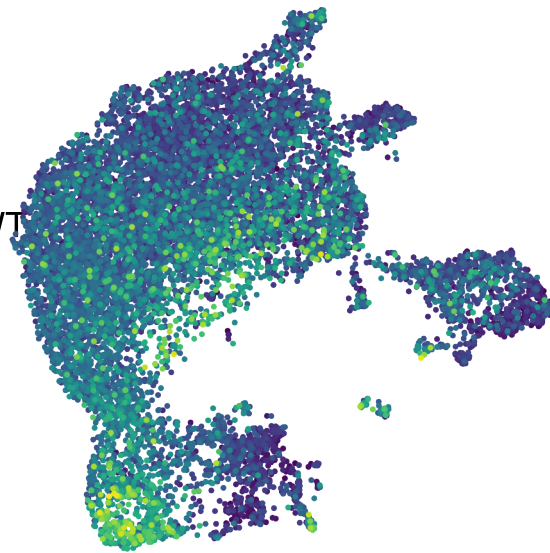


batch



total_counts

● Corey_WT
● Yi_WT
● Yi_KO



n_genes_by_counts

