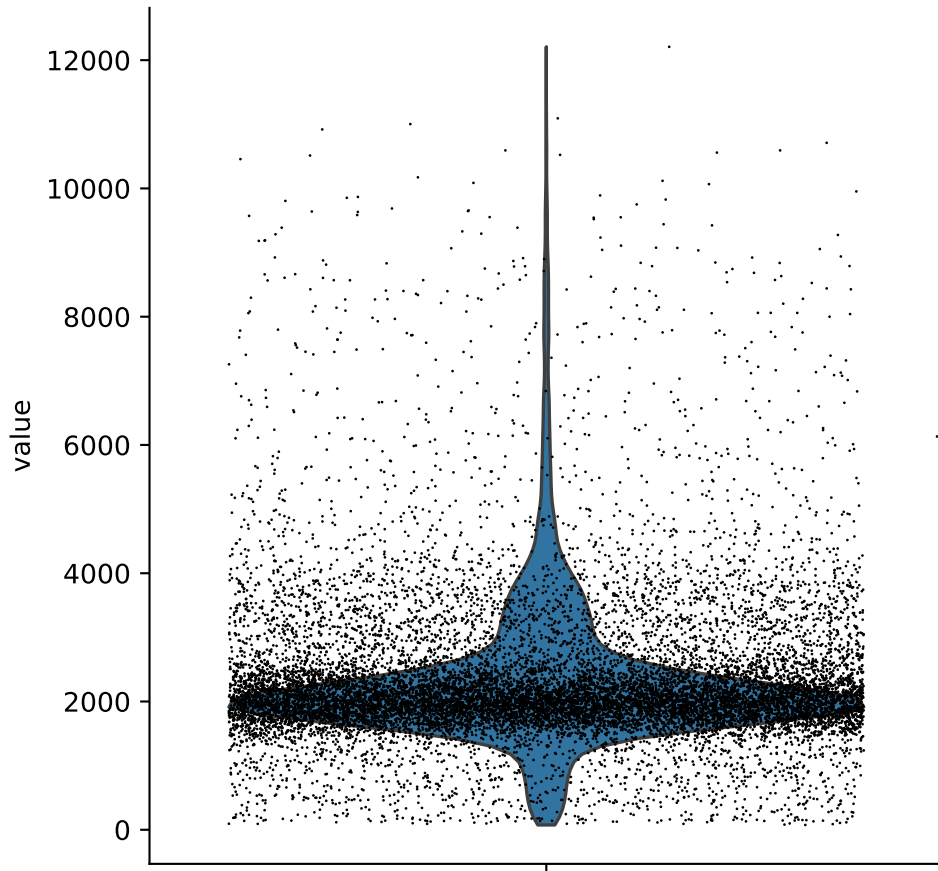
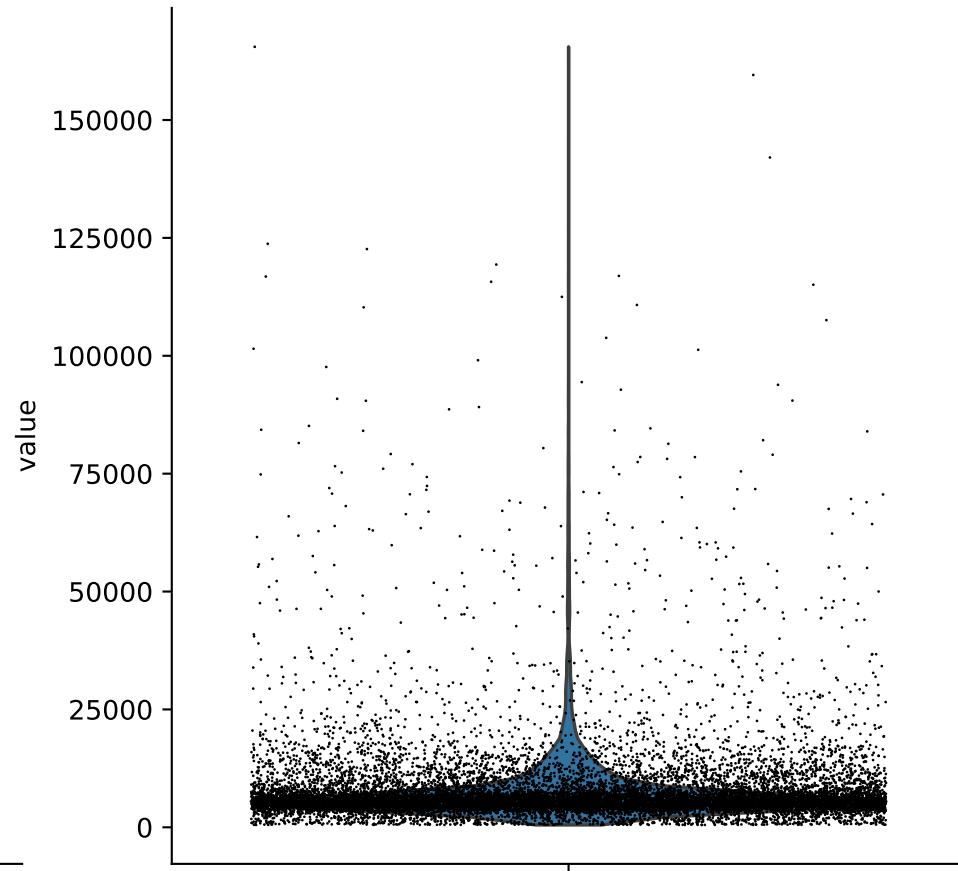


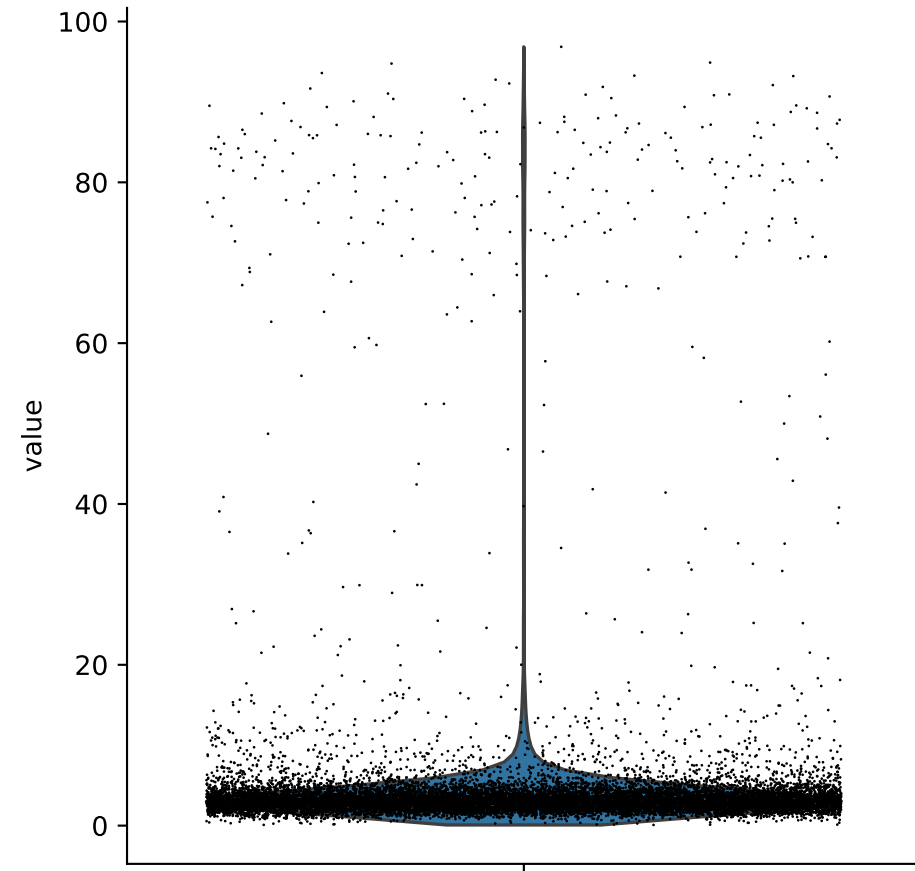
n_genes_by_counts



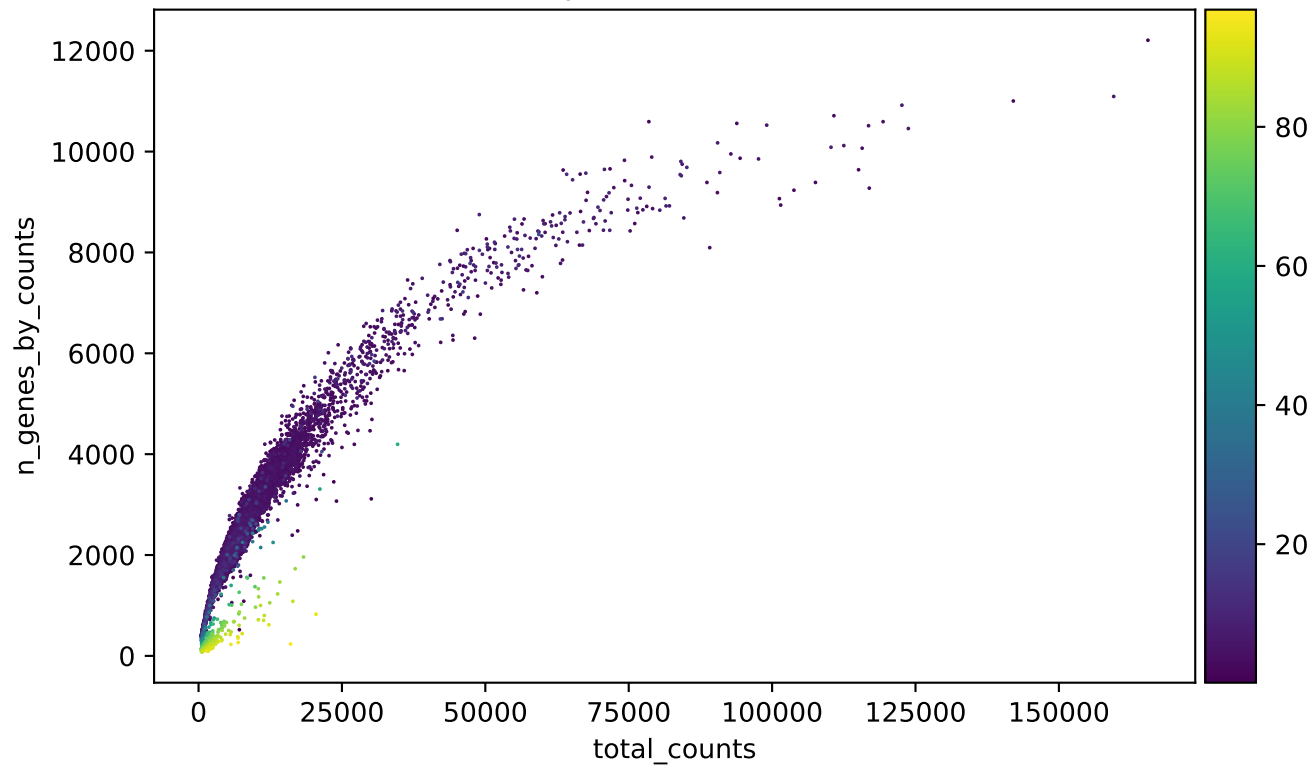
total_counts

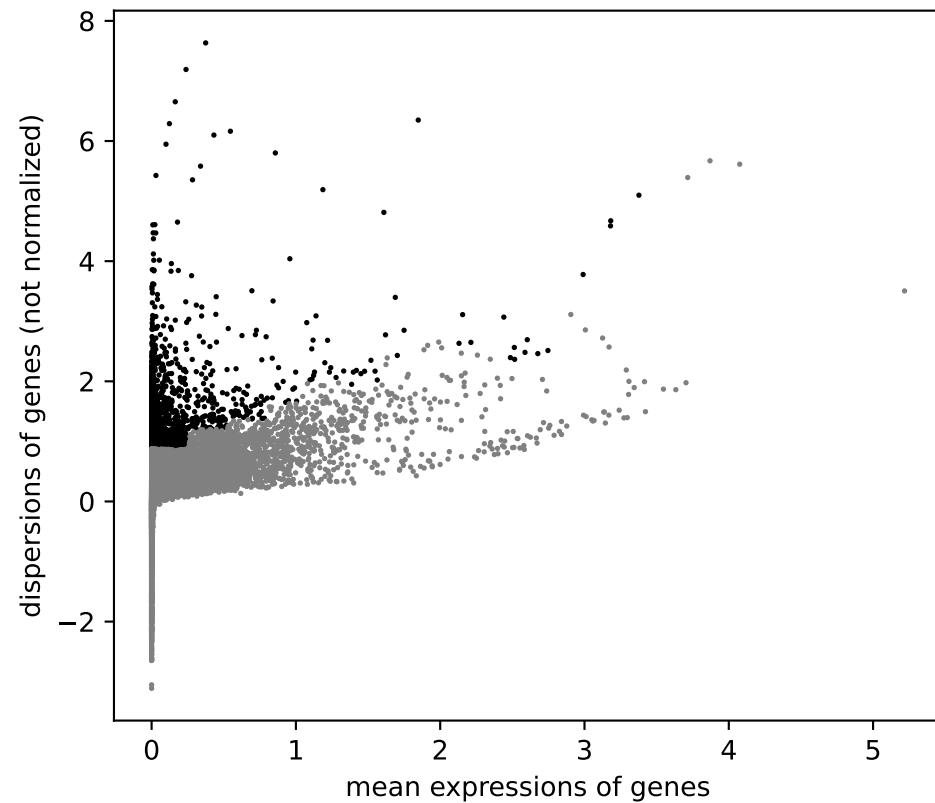
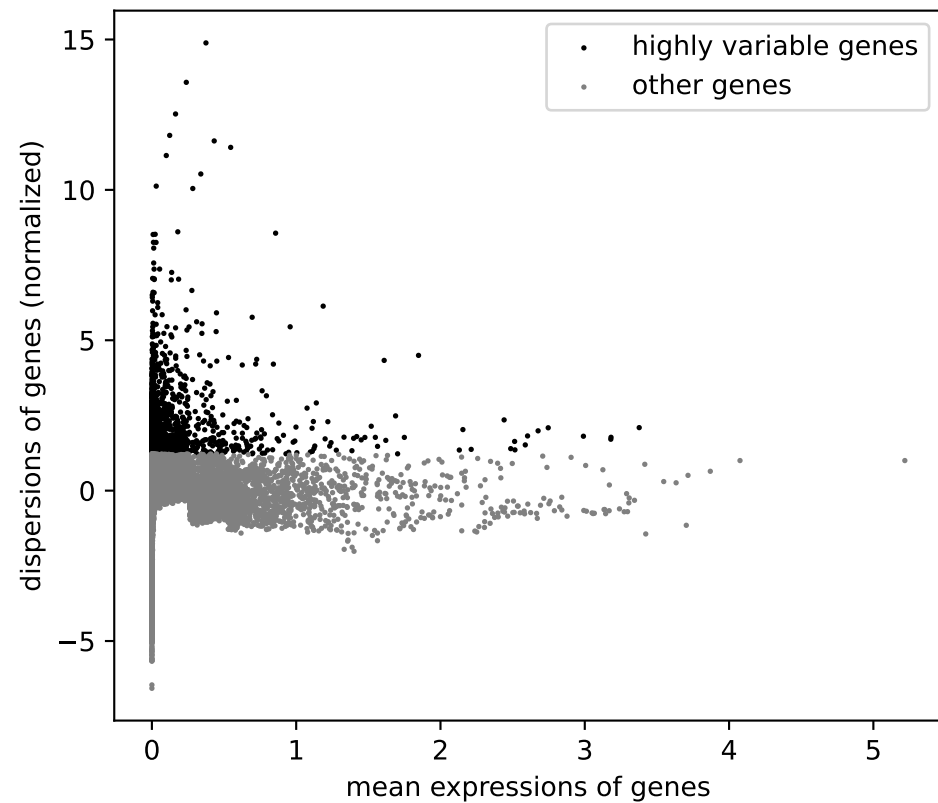


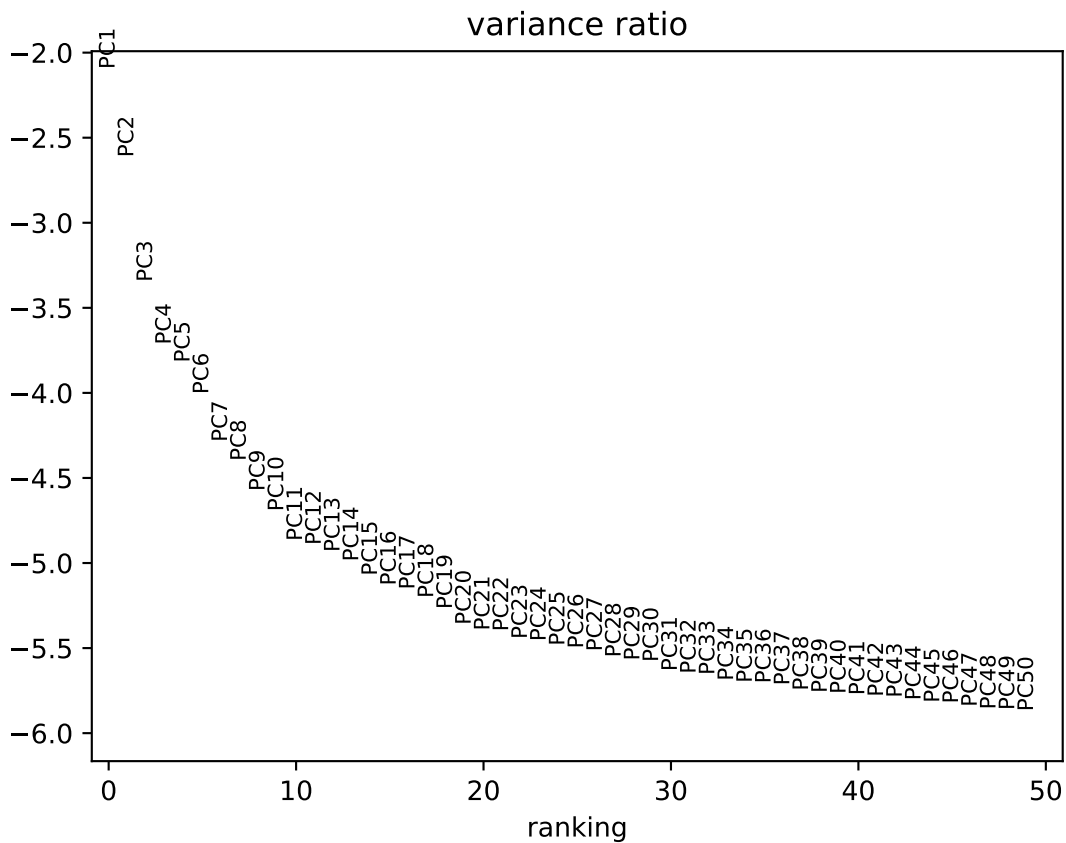
pct_counts_mt

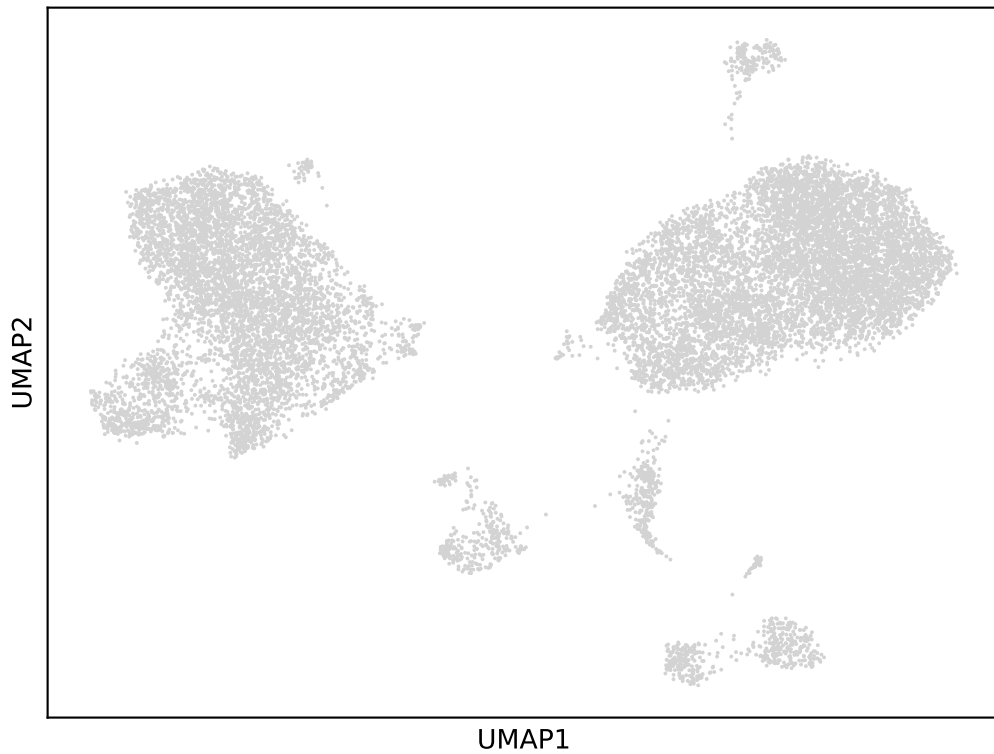


pct counts mt

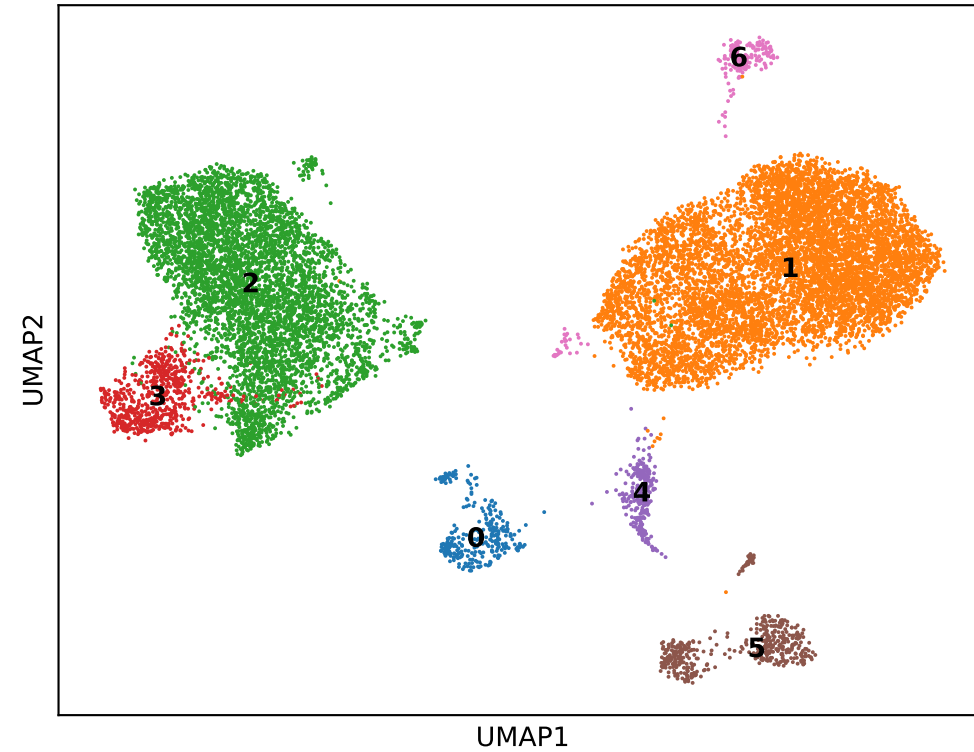




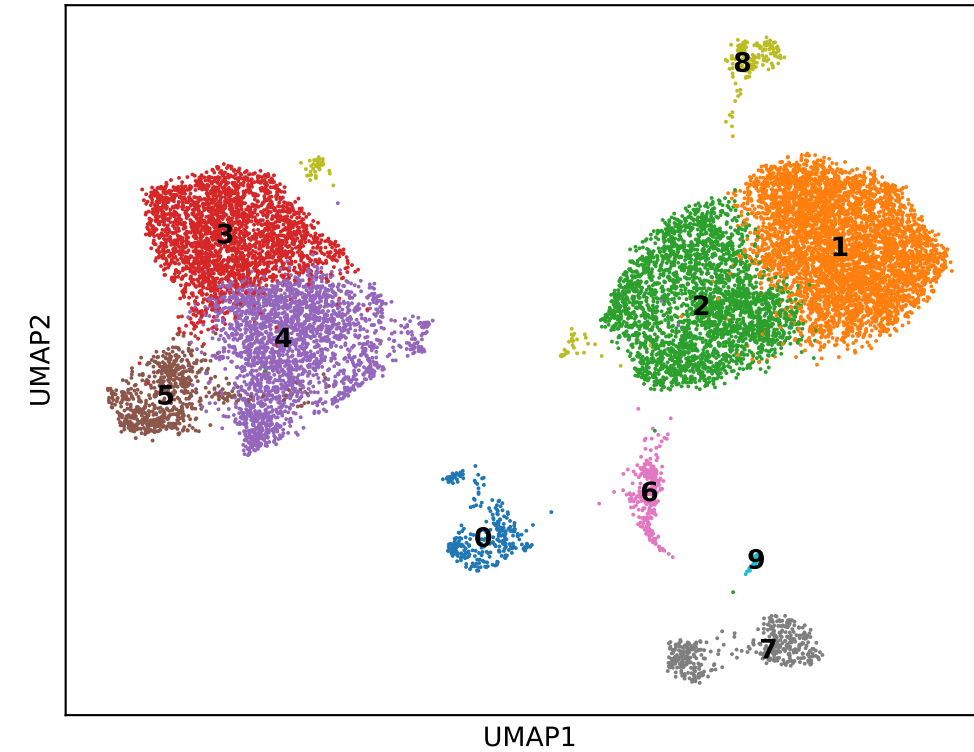




leiden_res_0.20



leiden_res_0.50



leiden_res_0.80

