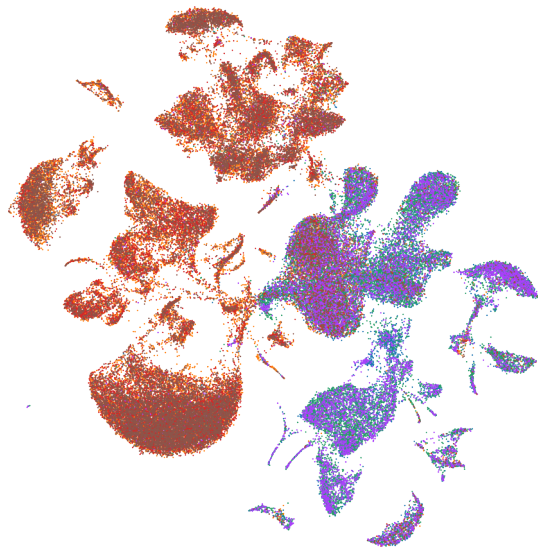
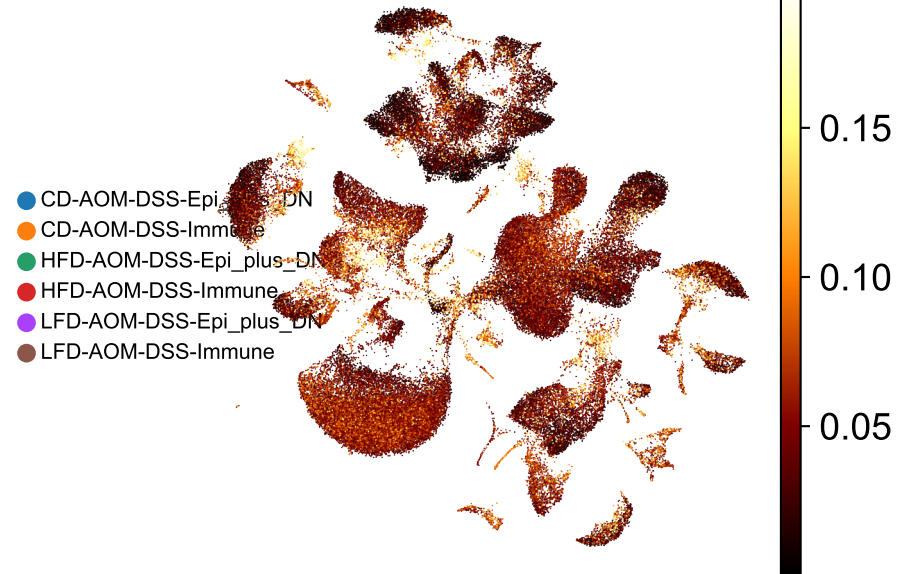


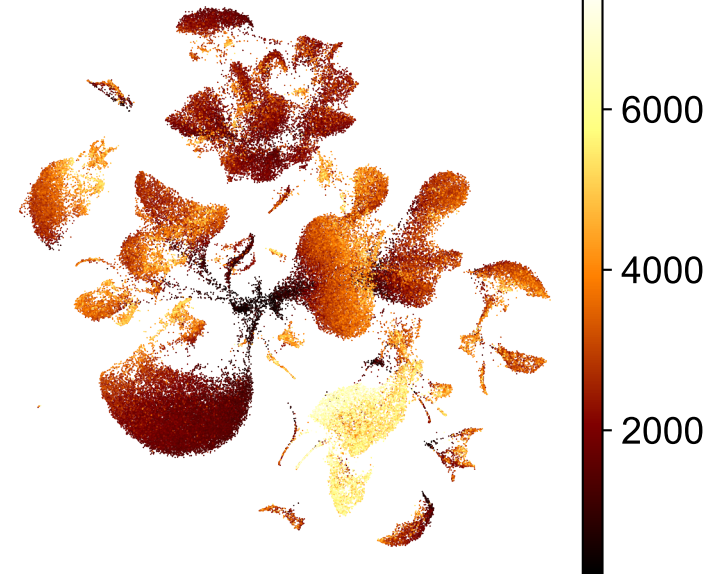
condition



doublet_score



n_genes_by_counts



pct_counts_mt

