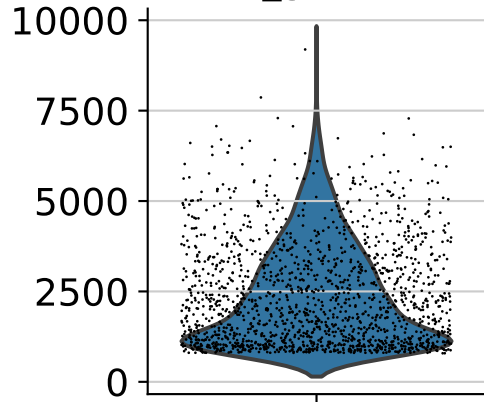
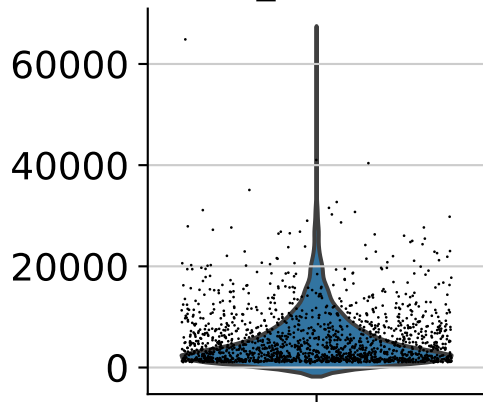


n_genes



n_counts



percent_mito

