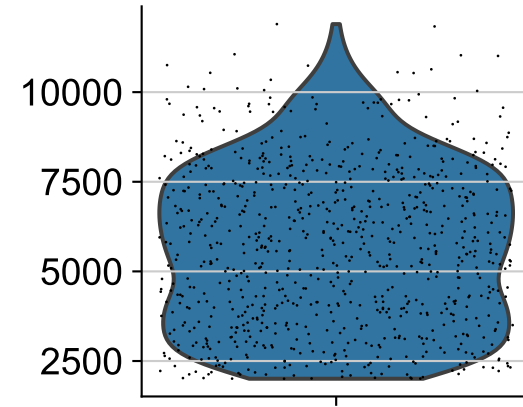
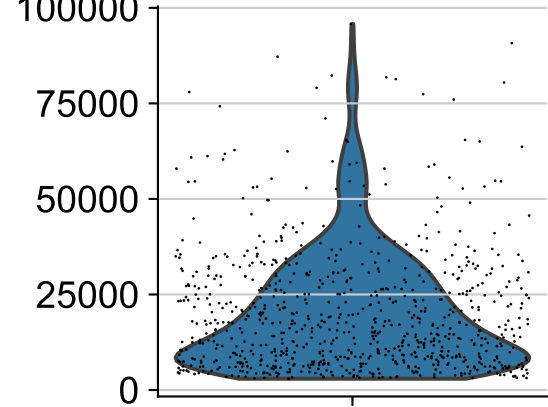


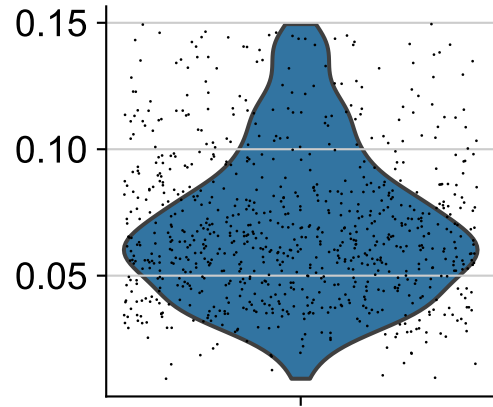
n_genes



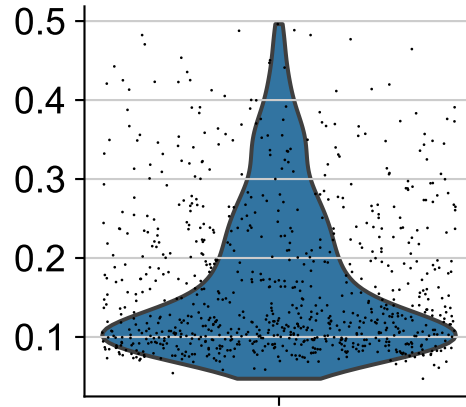
n_counts



percent_mito



ERCC_genes



percent_ribo

