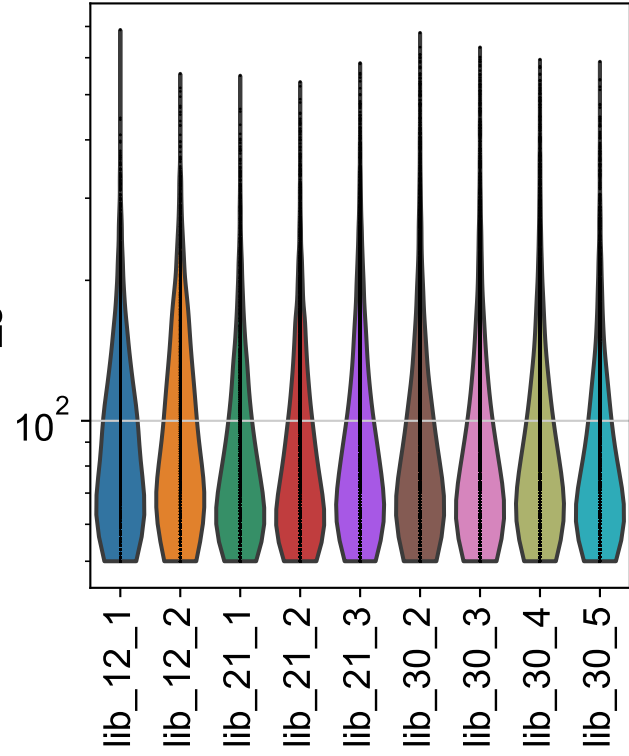
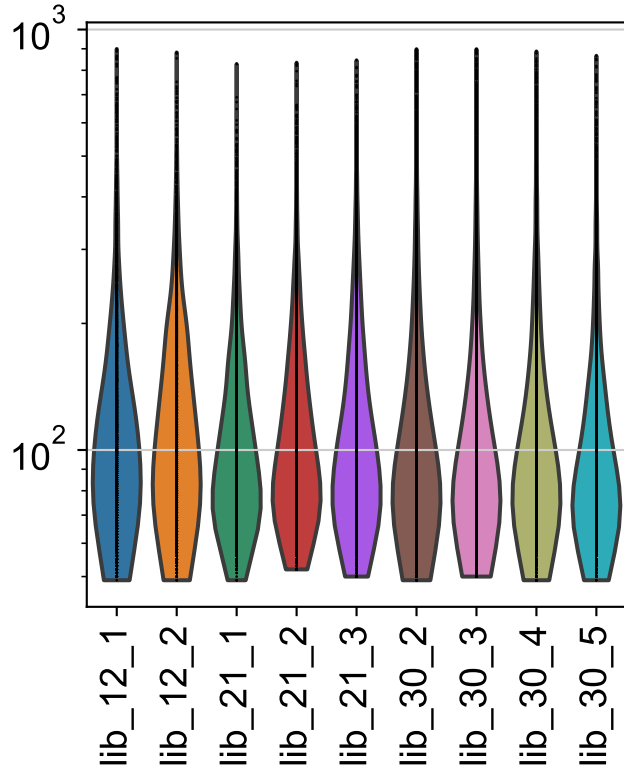


n_genes



total_counts



n_counts

