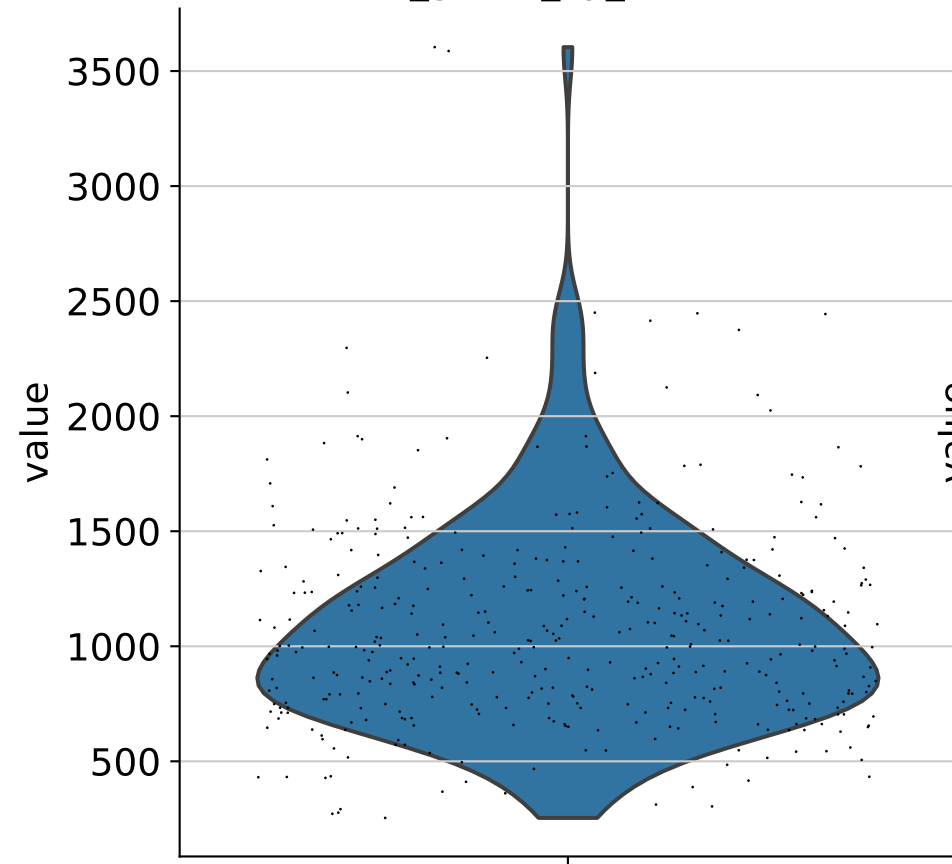
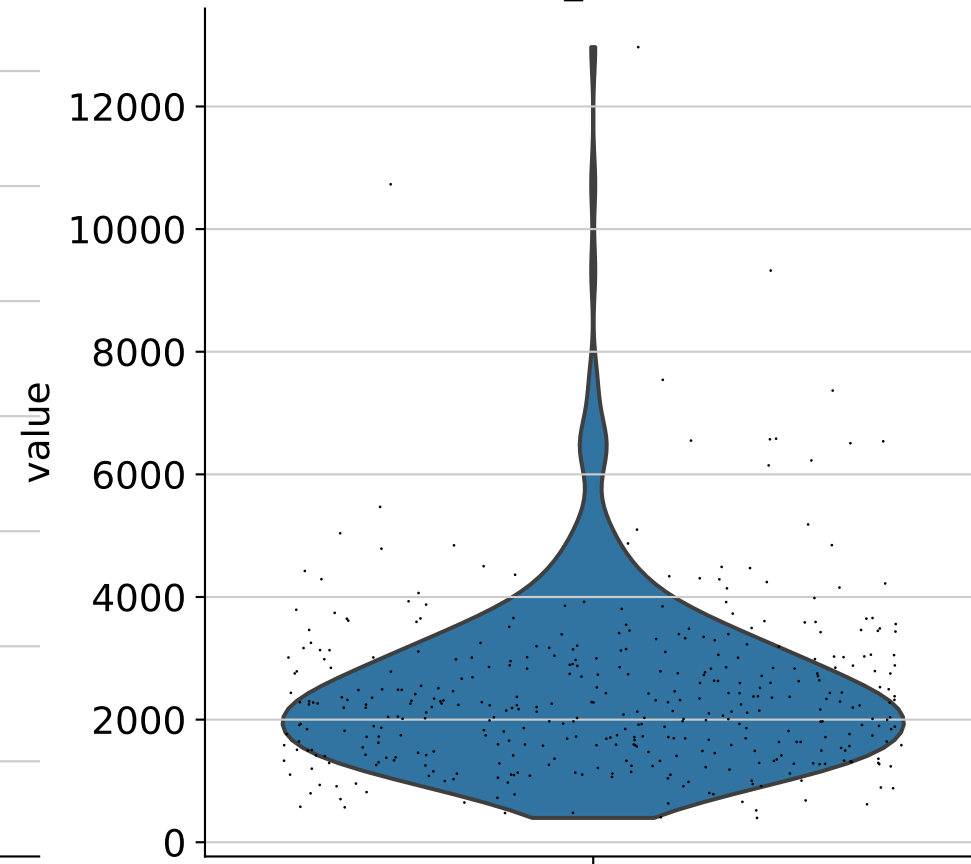


n_genes_by_counts



total_counts



pct_counts_mt

