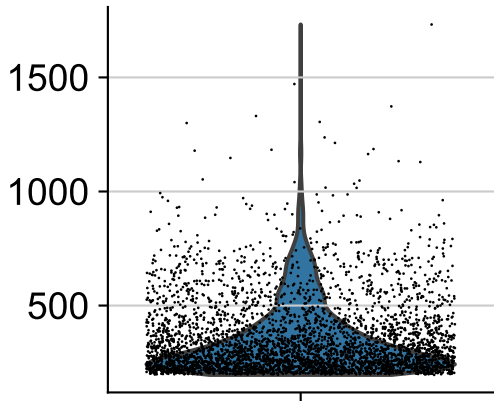


n_genes_by_counts



total_counts

