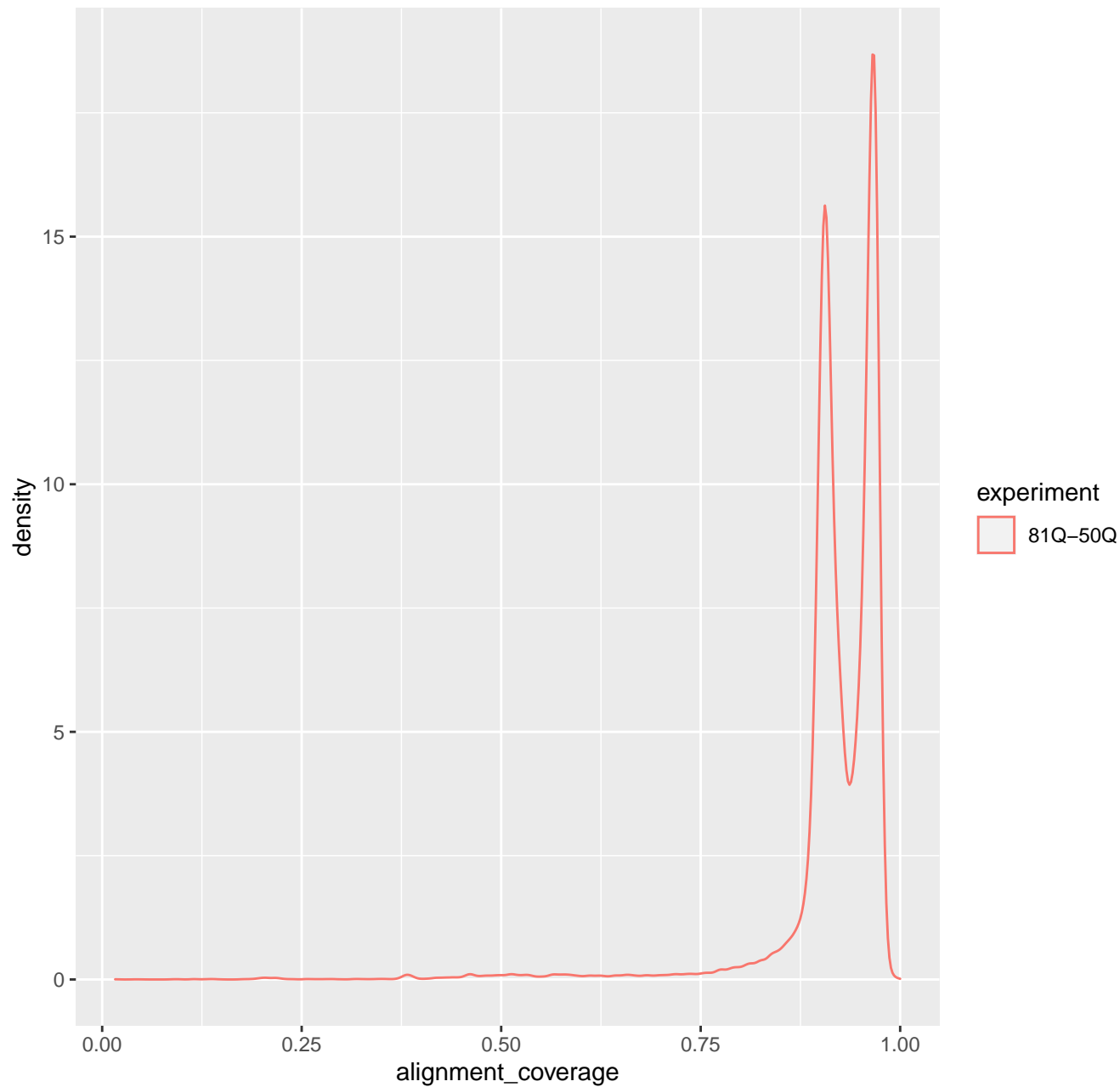
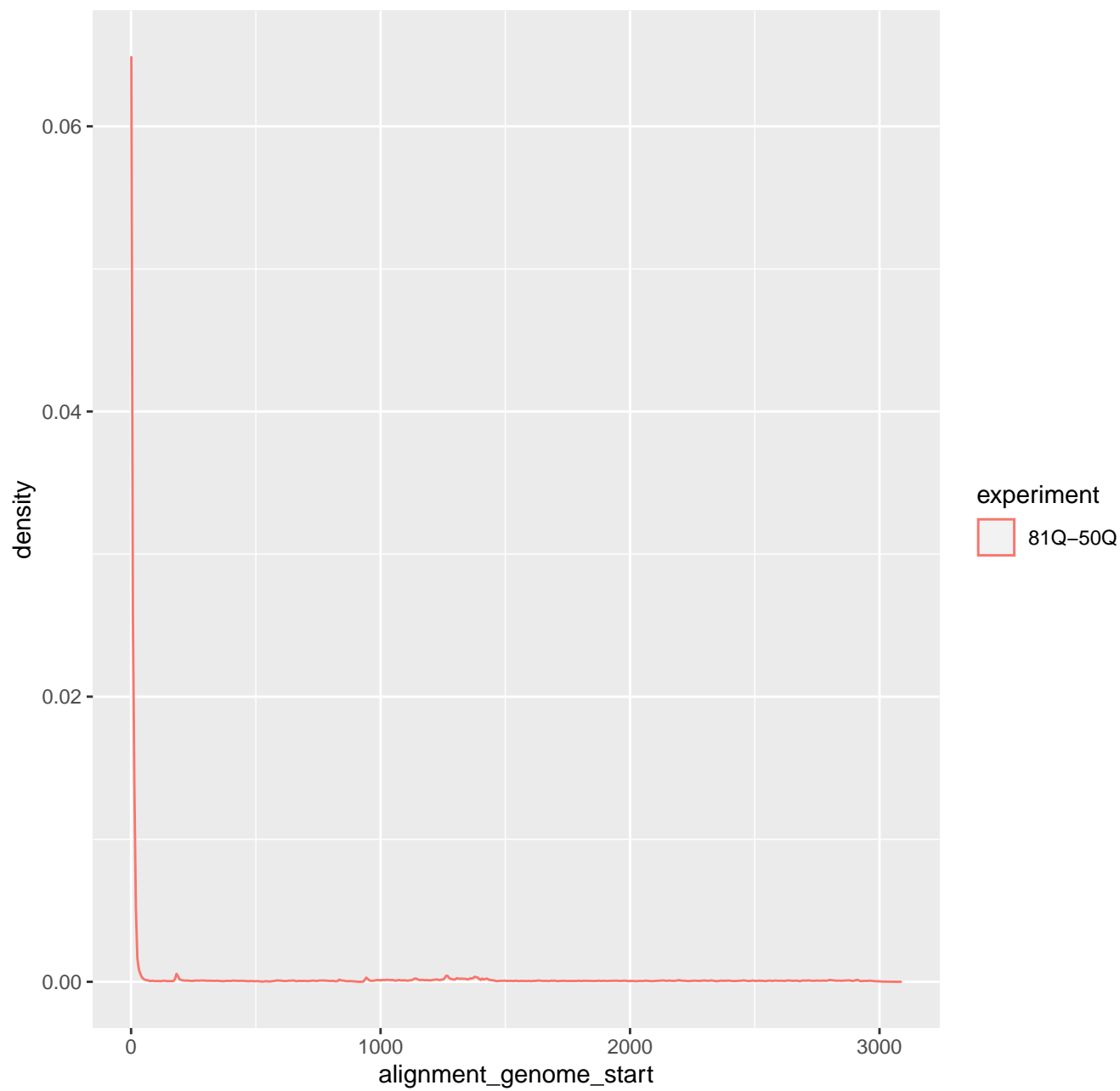


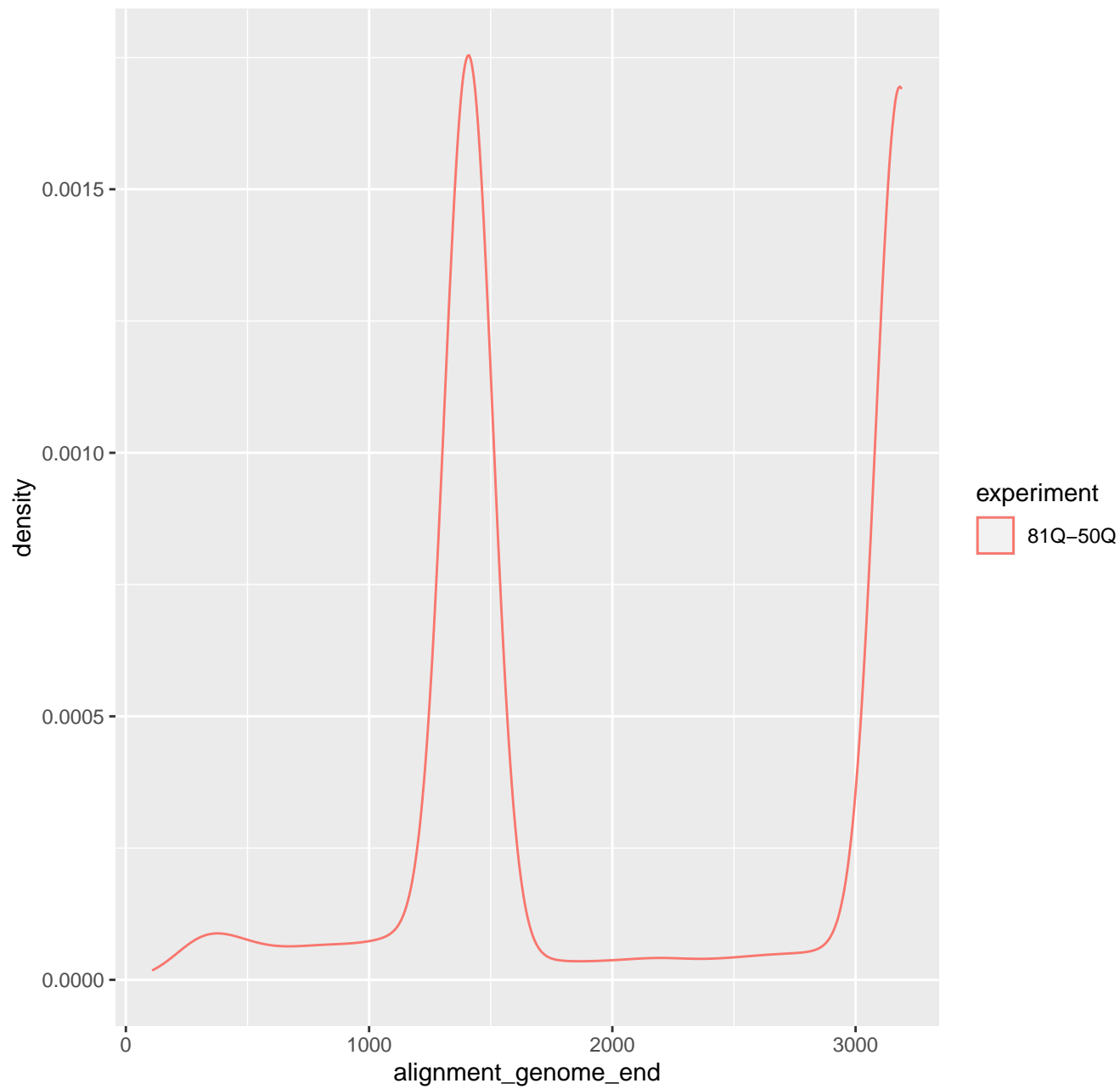
# Alignment Coverage



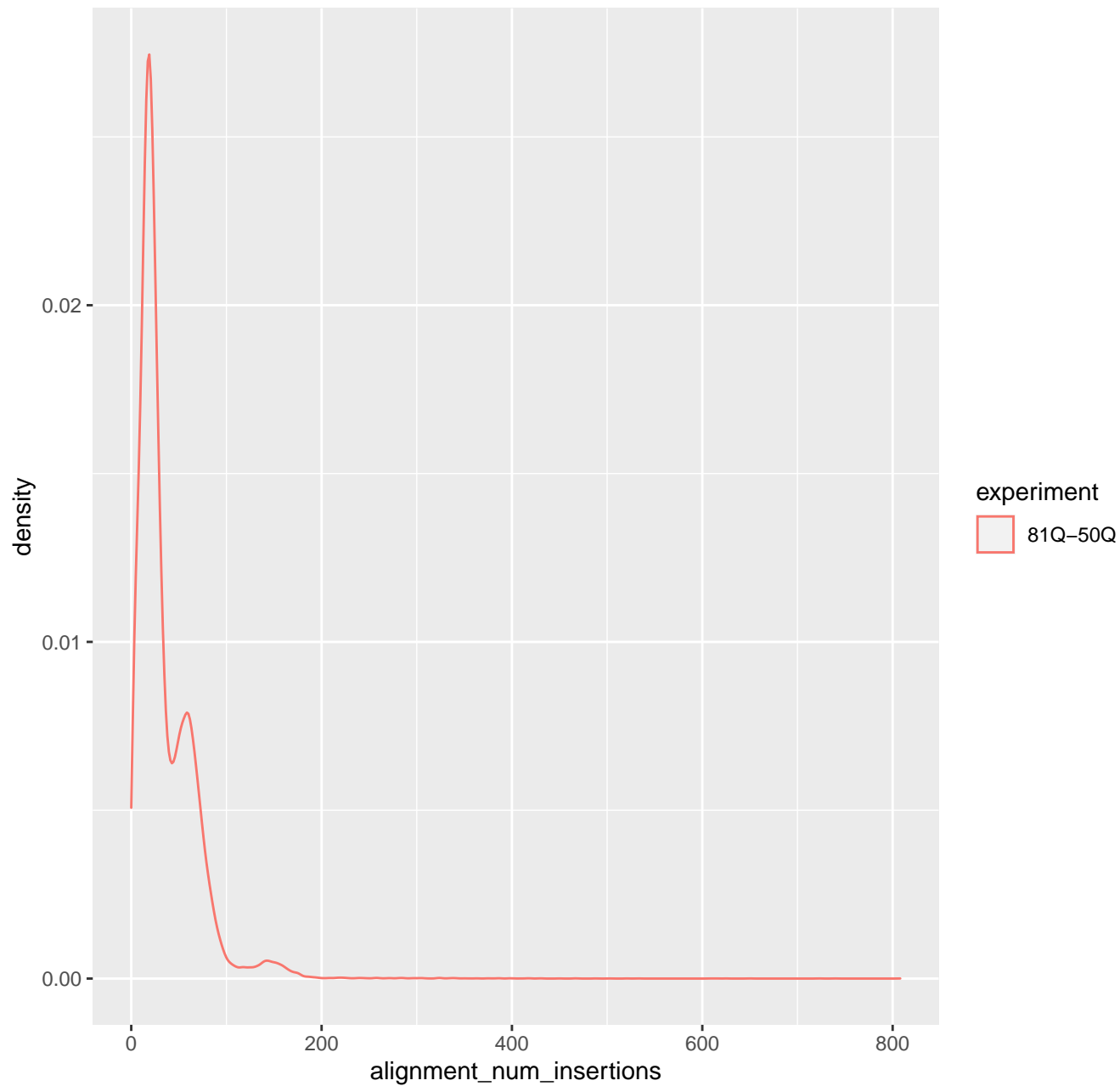
Alignment genome\_start



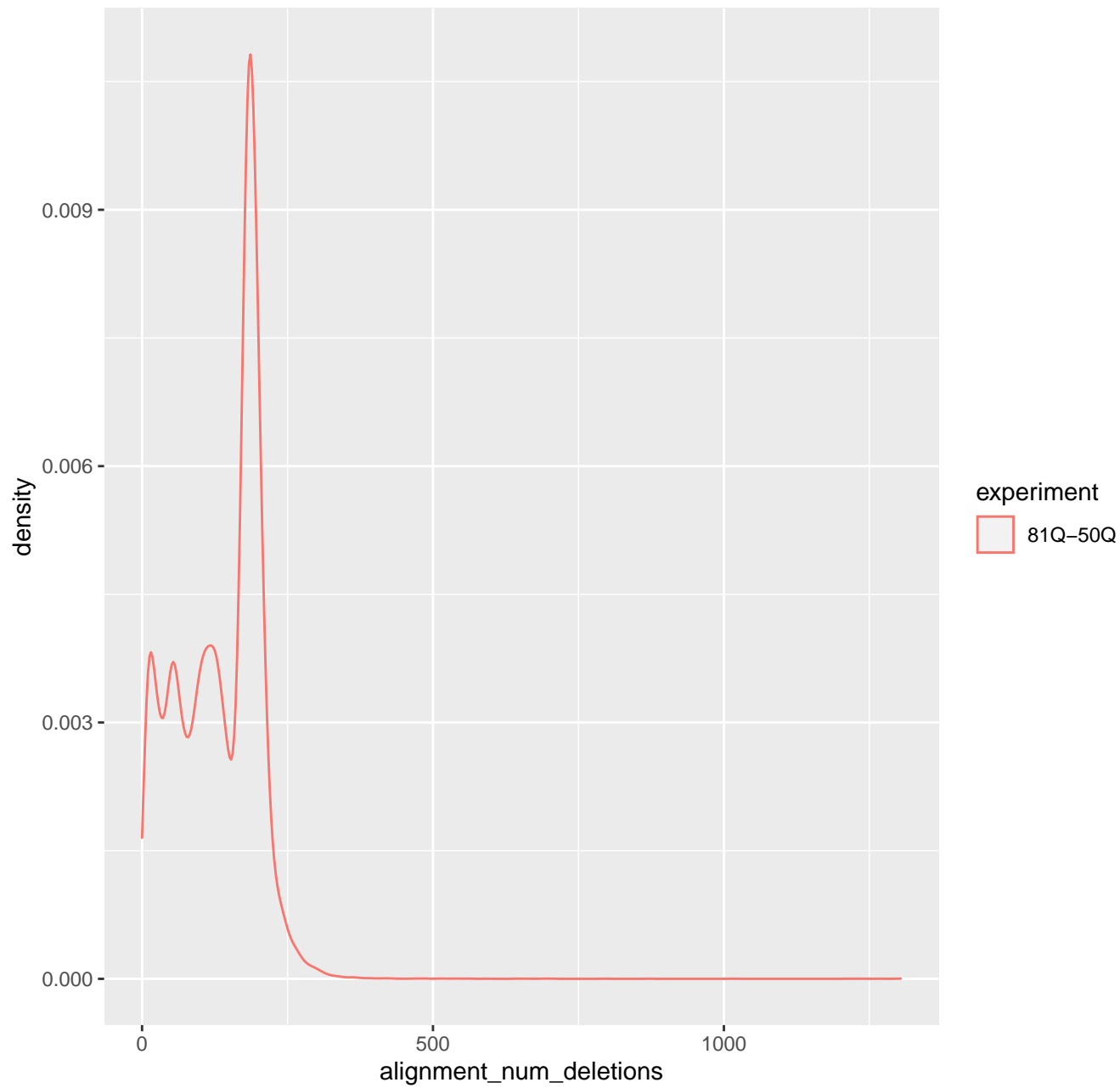
Alignment genome\_end



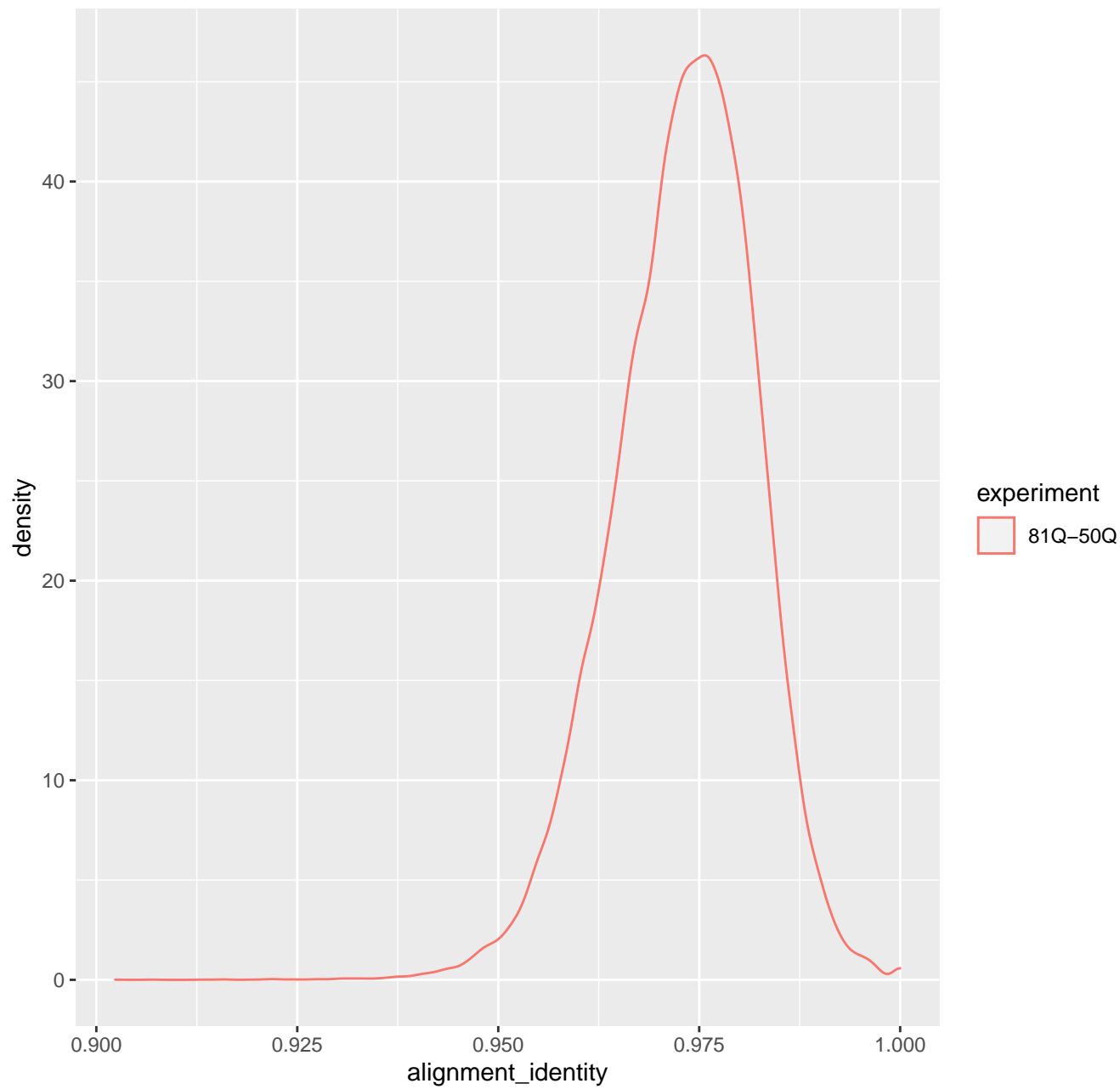
Number of insertions



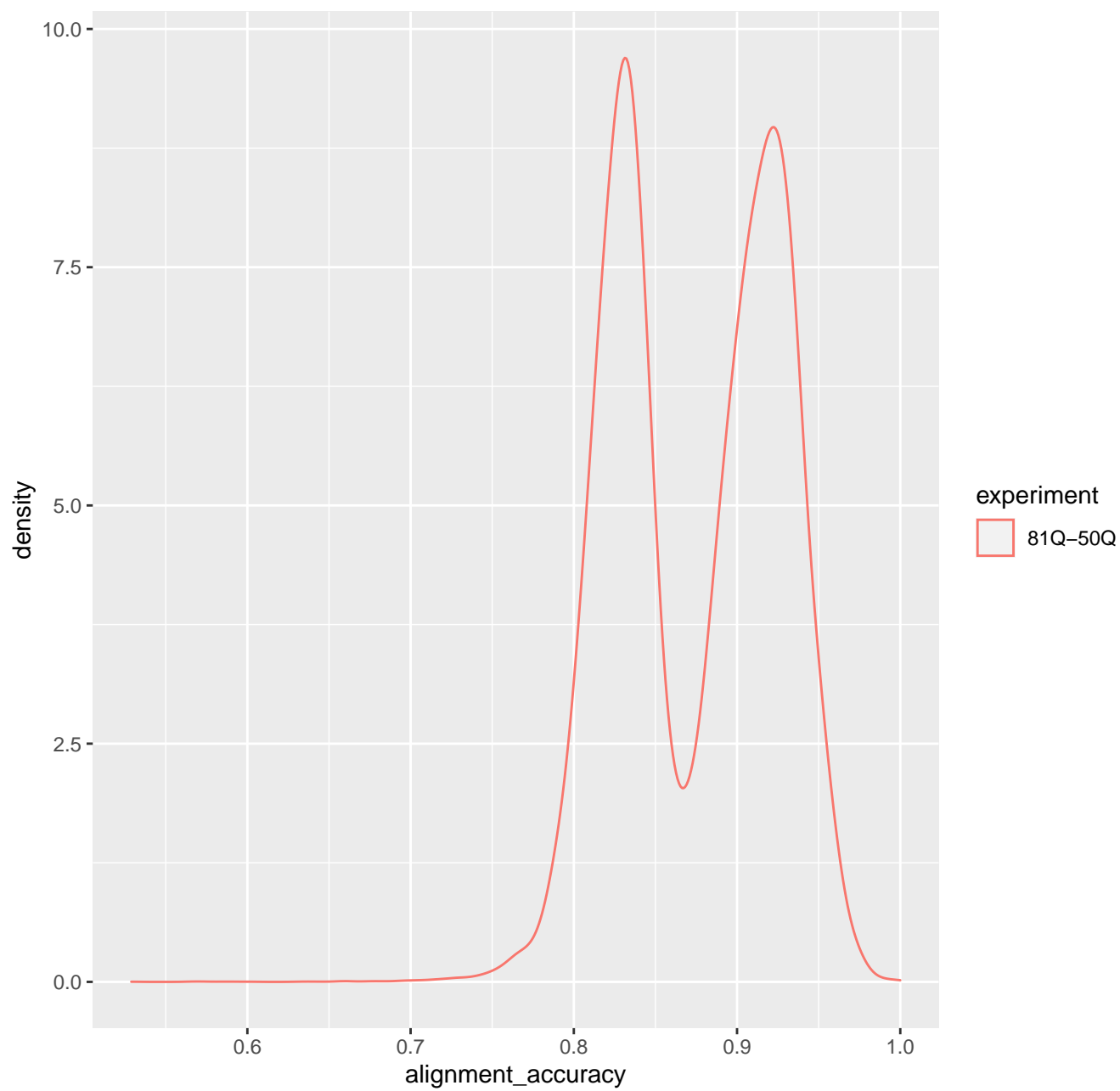
Number of deletions



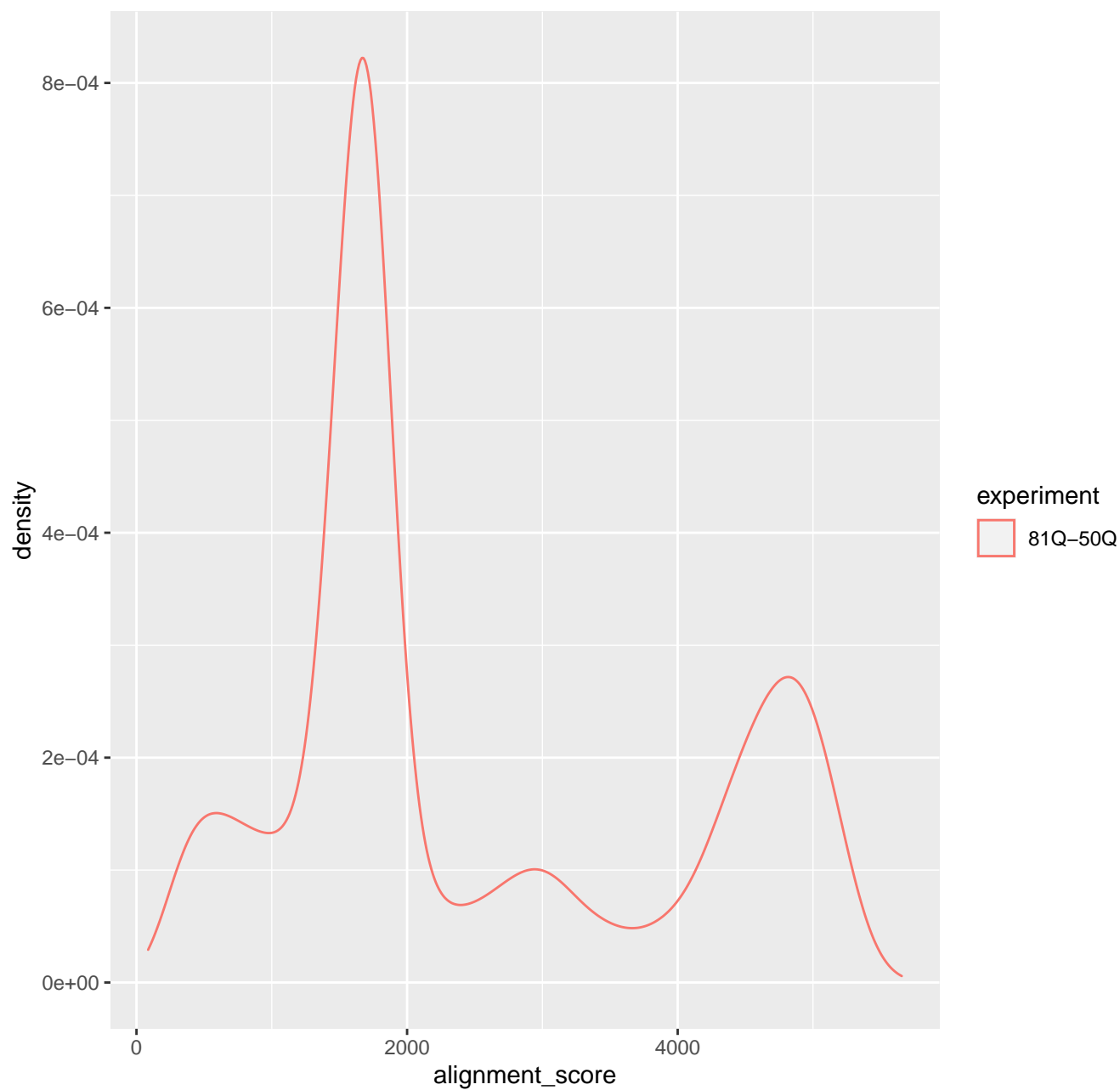
Alignment\_identity



Alignment\_accuracy

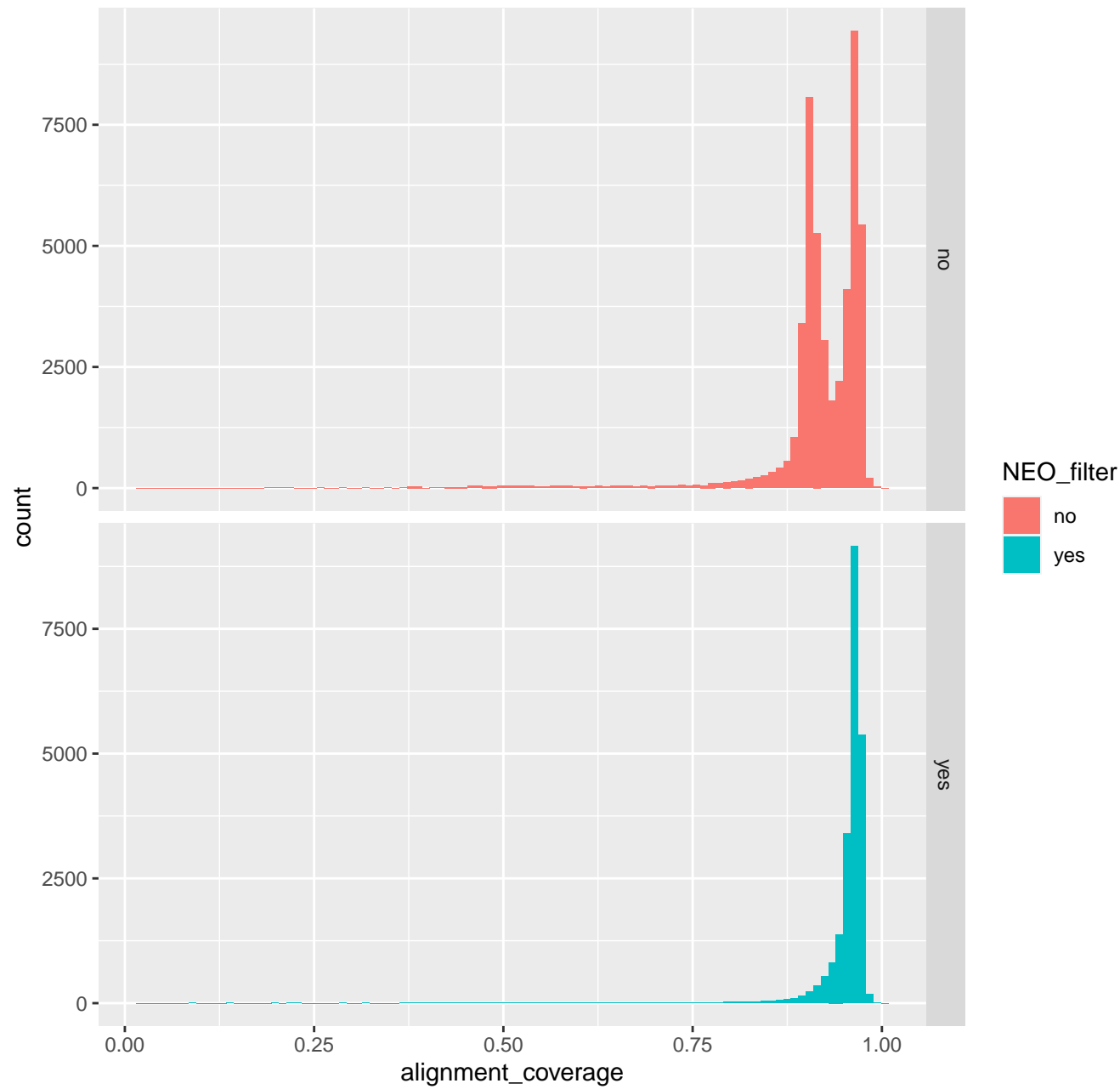


Alignment\_score

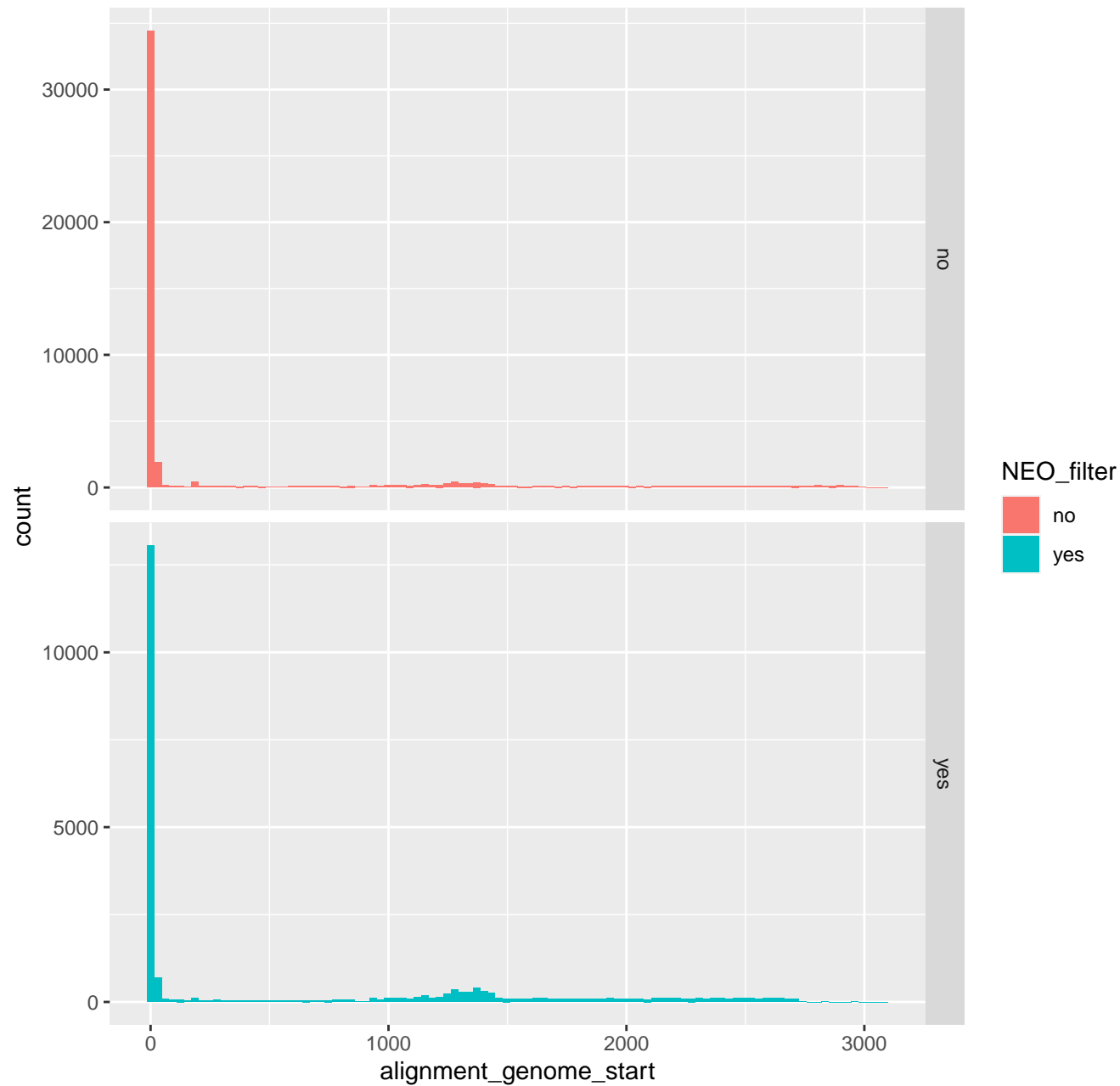




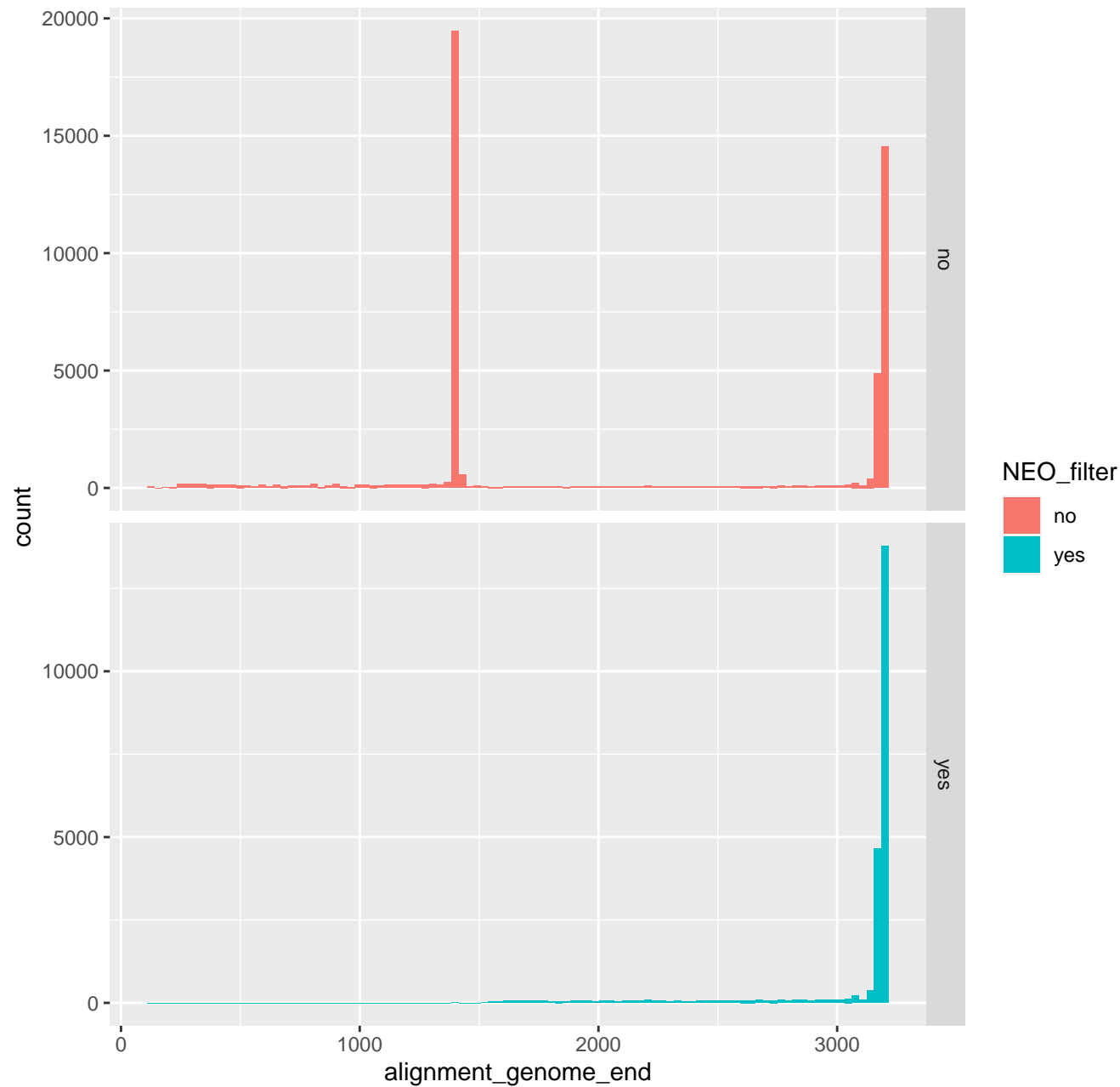
Alignment Coverage



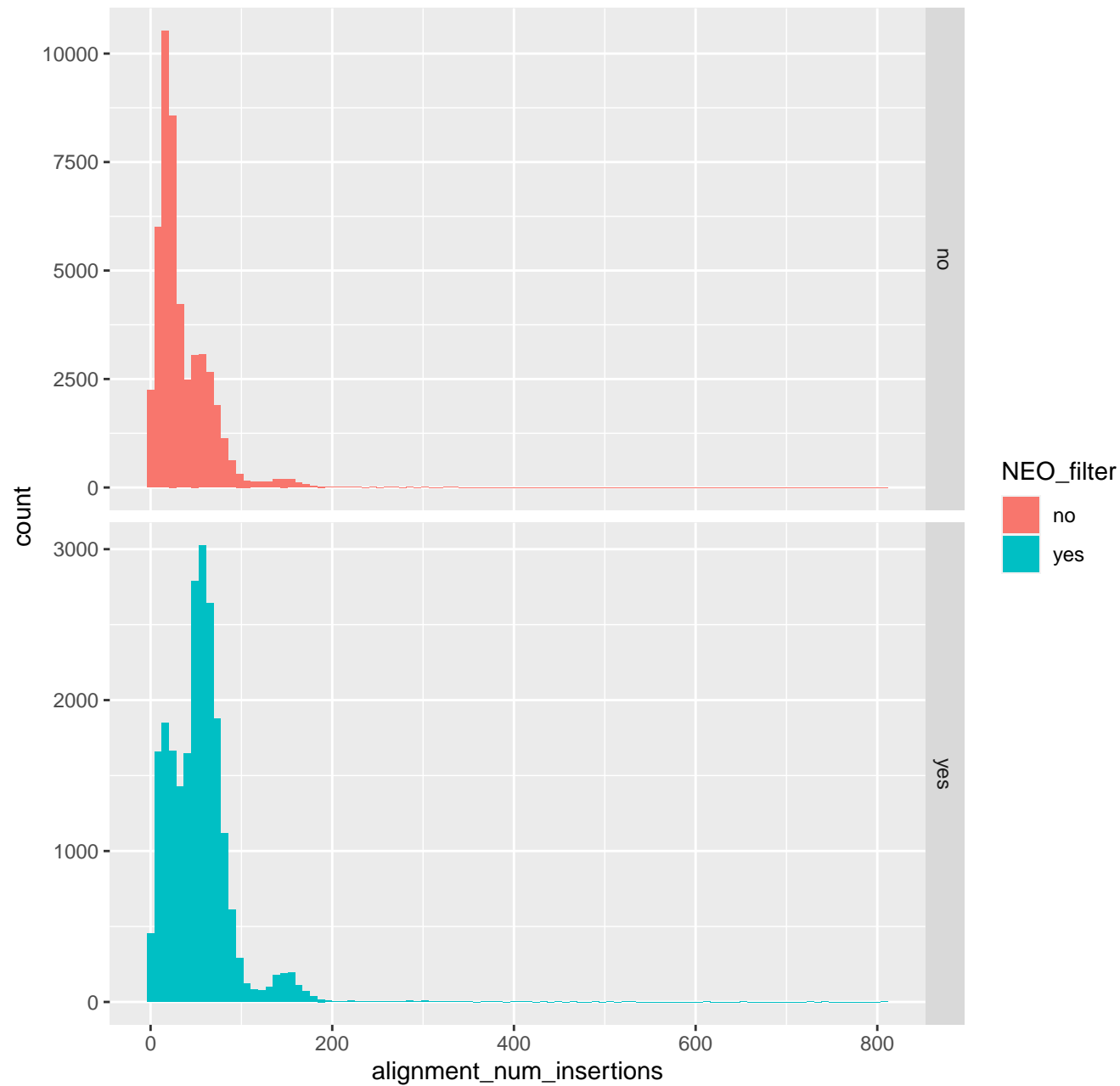
Alignment genome\_start



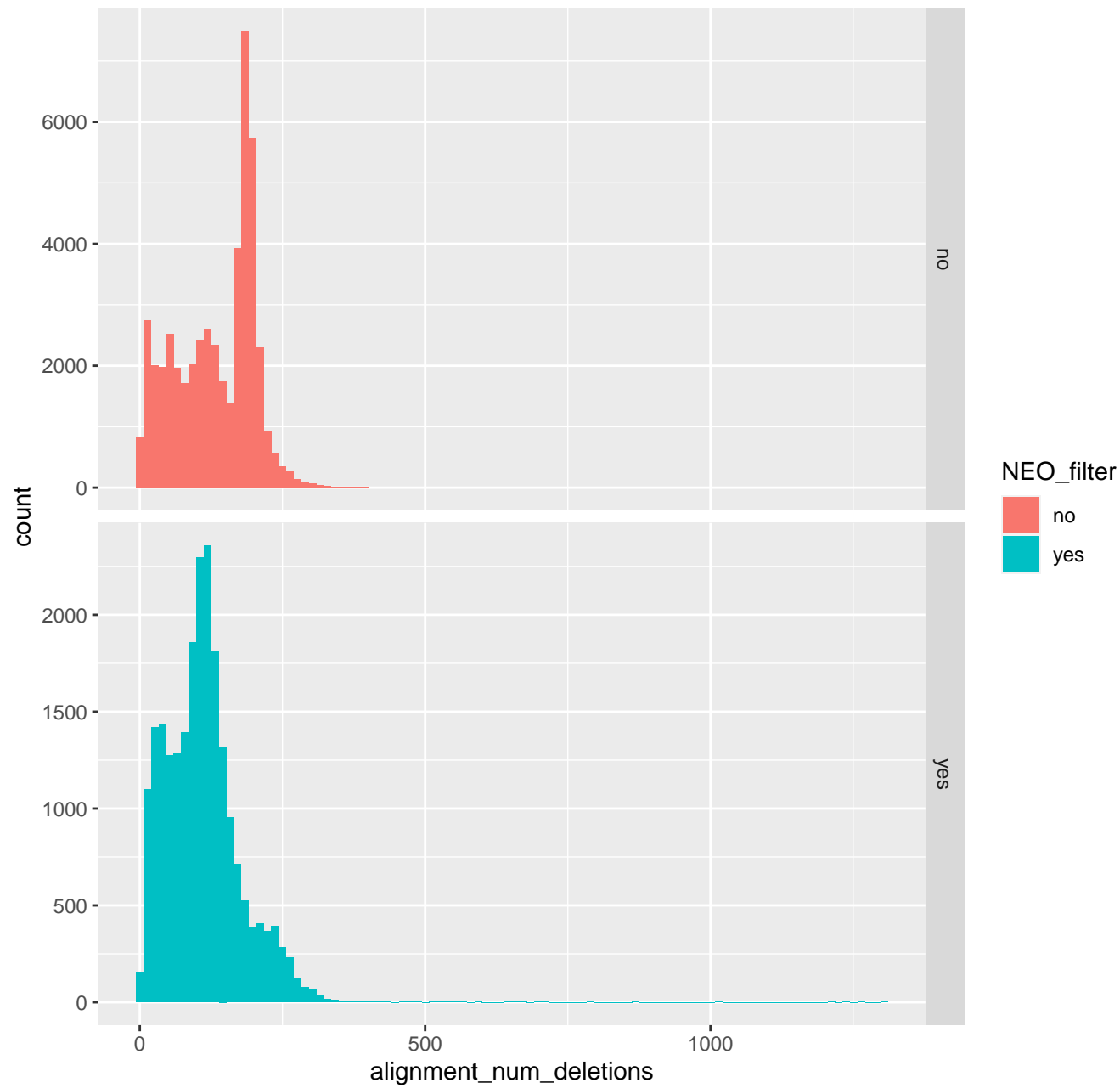
Alignment genome\_end



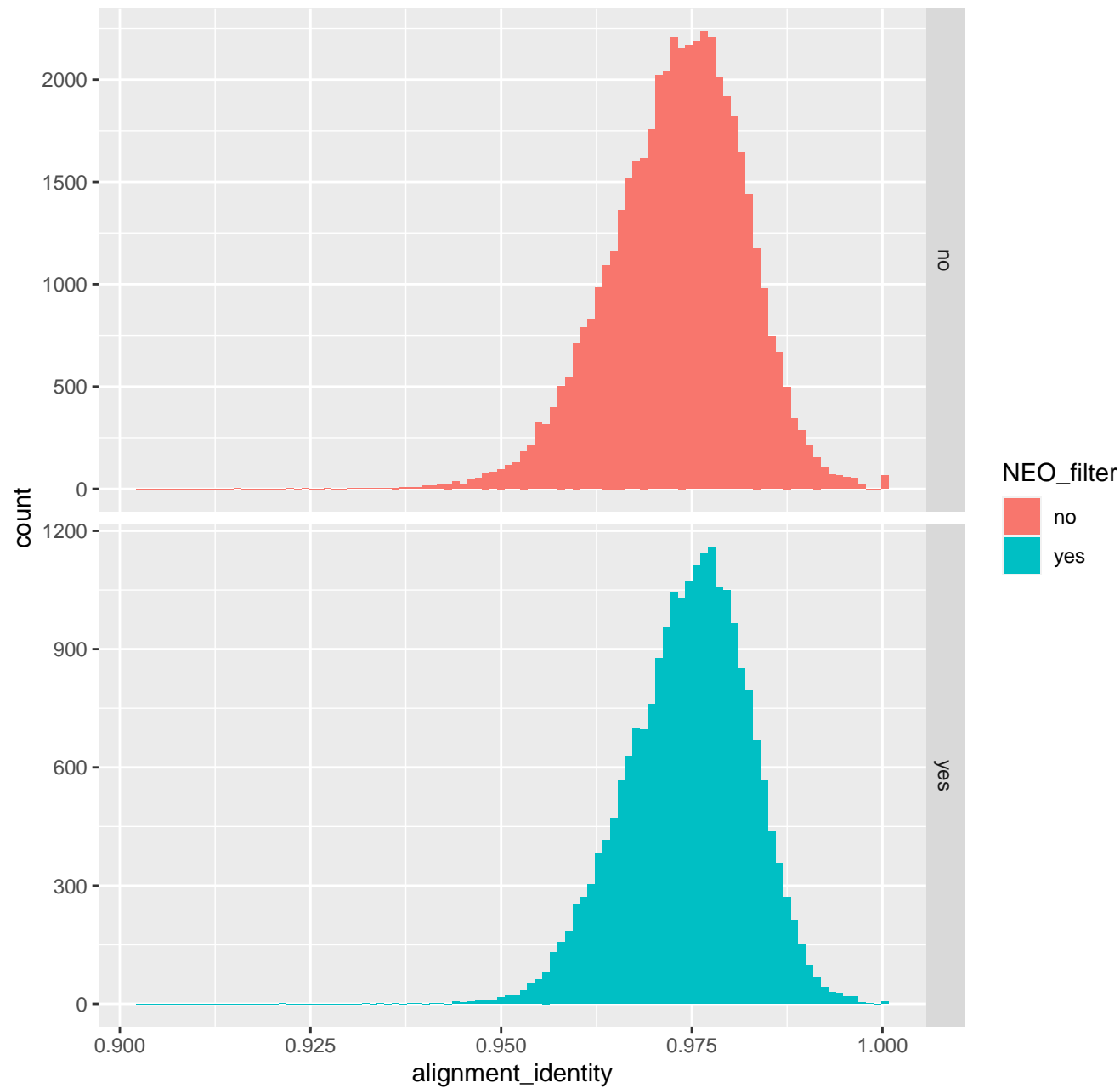
Number of insertions



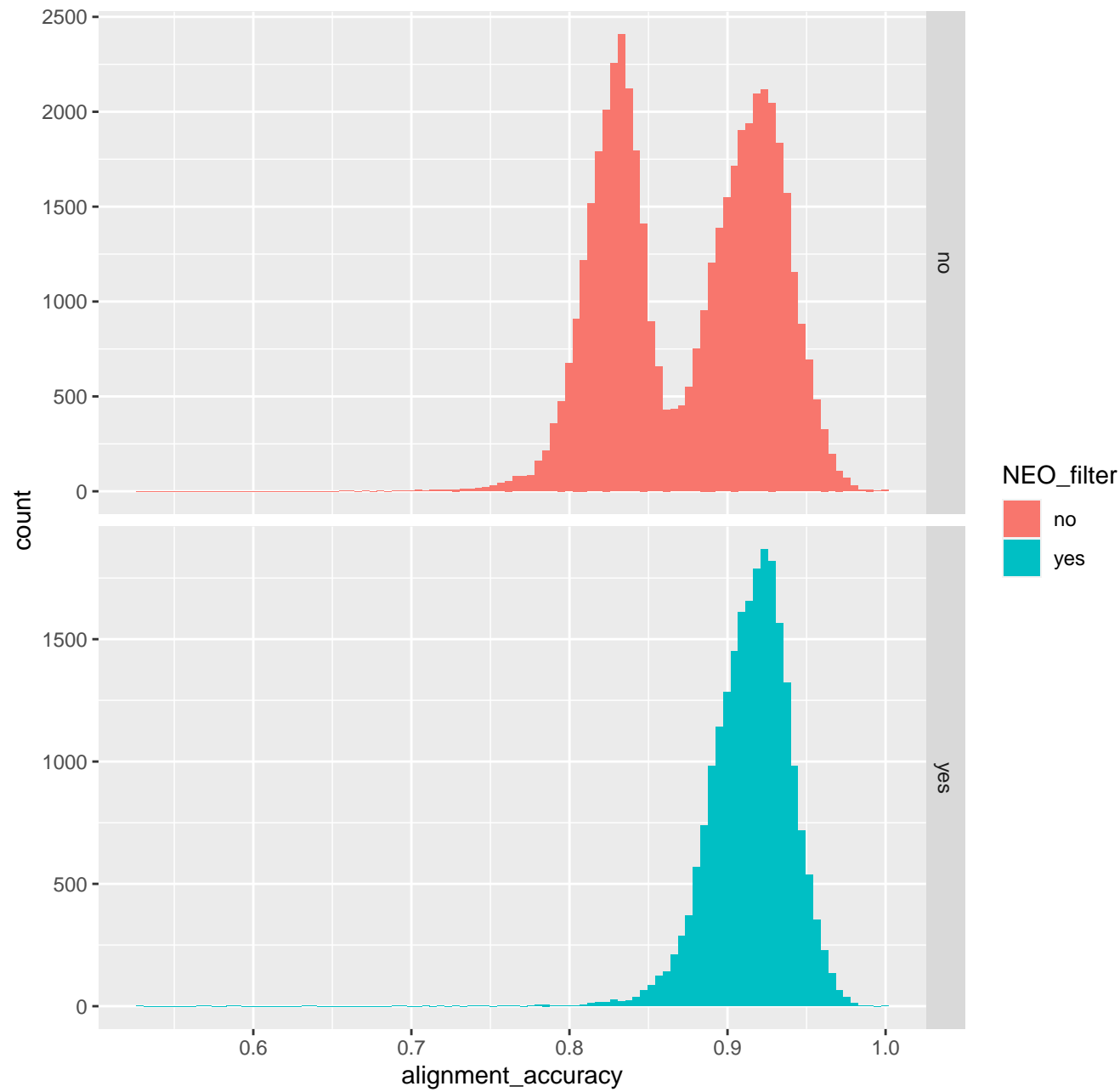
Number of deletions



Alignment\_identity



Alignment\_accuracy



Alignment\_score

