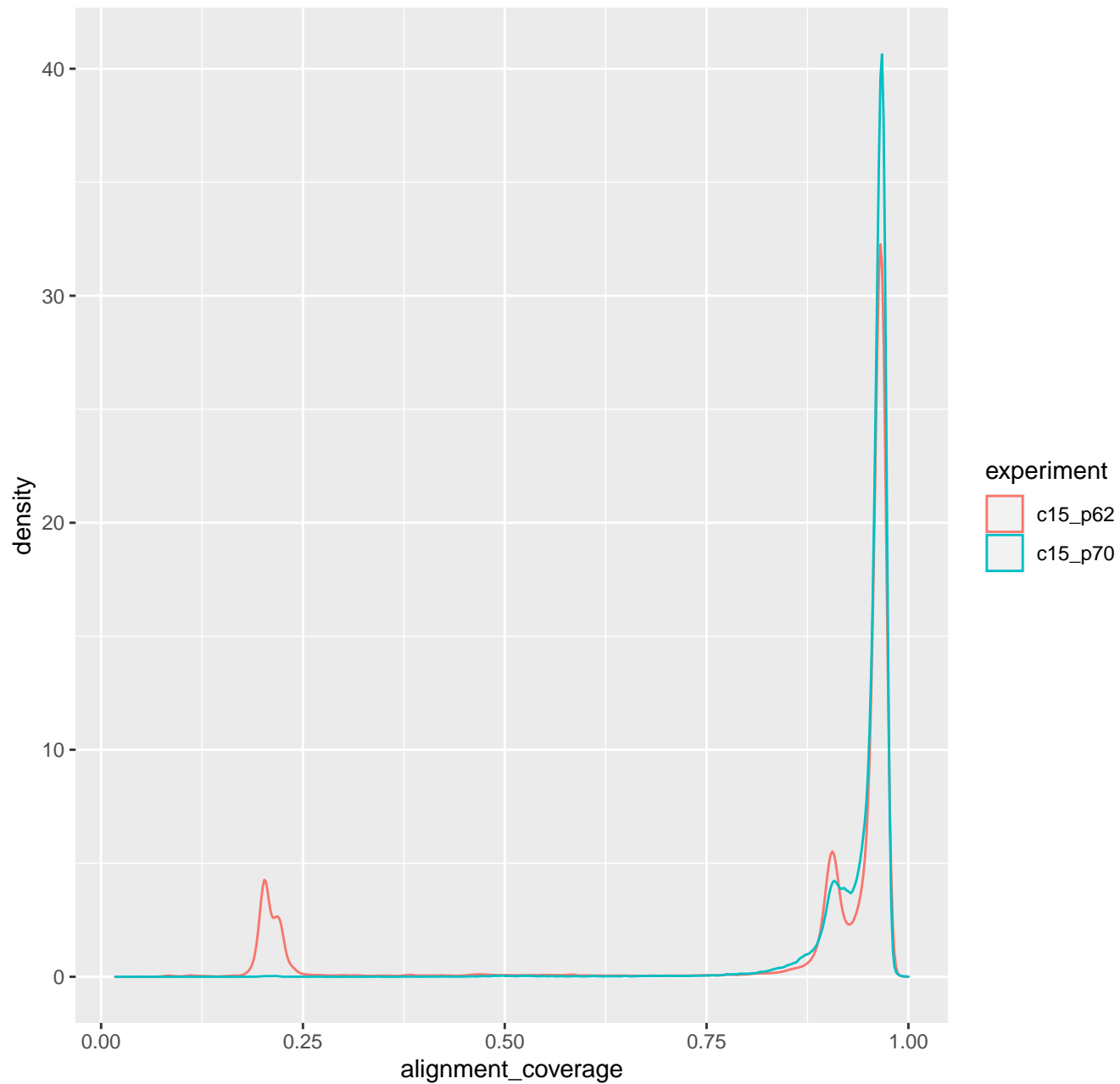
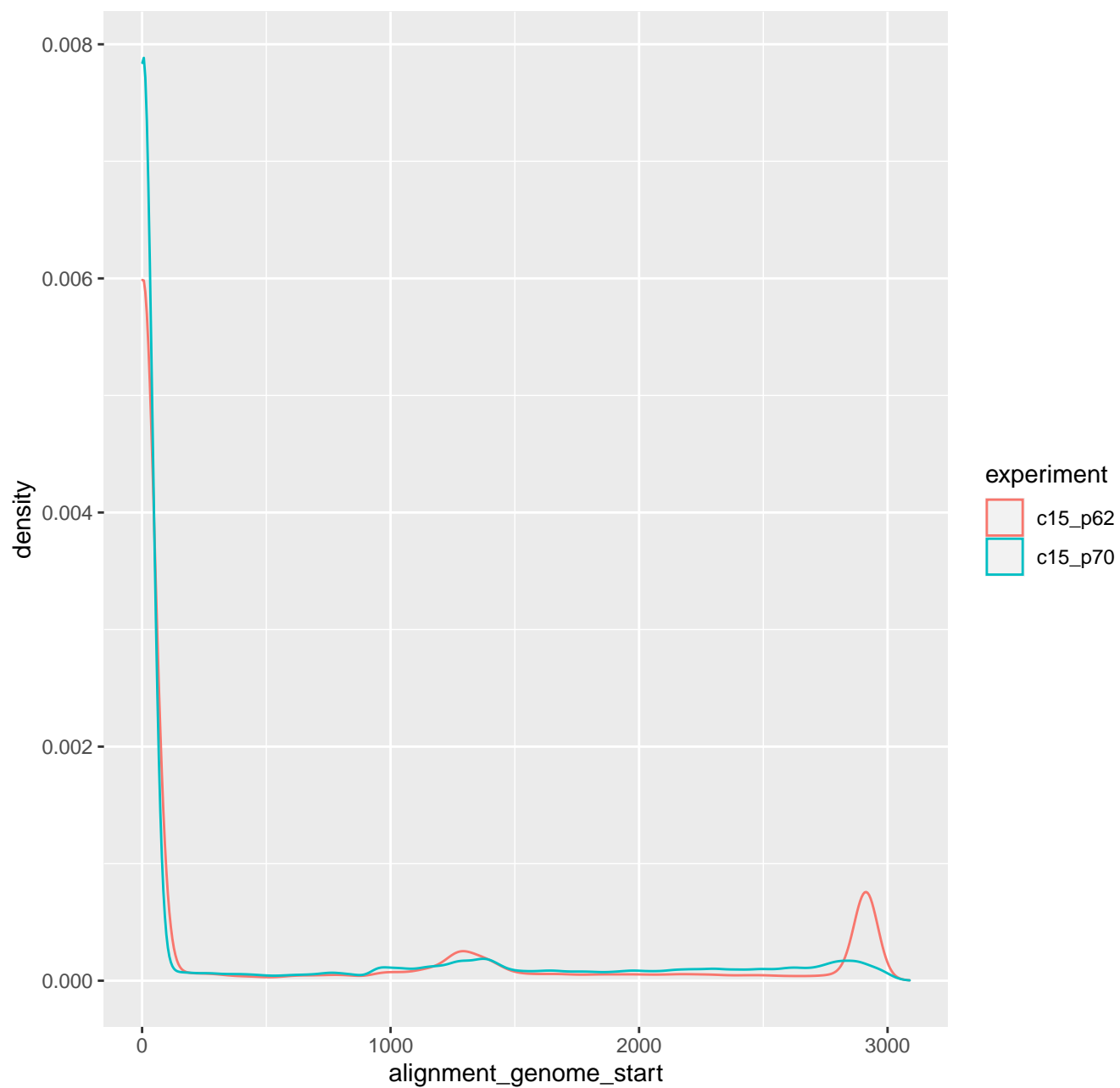


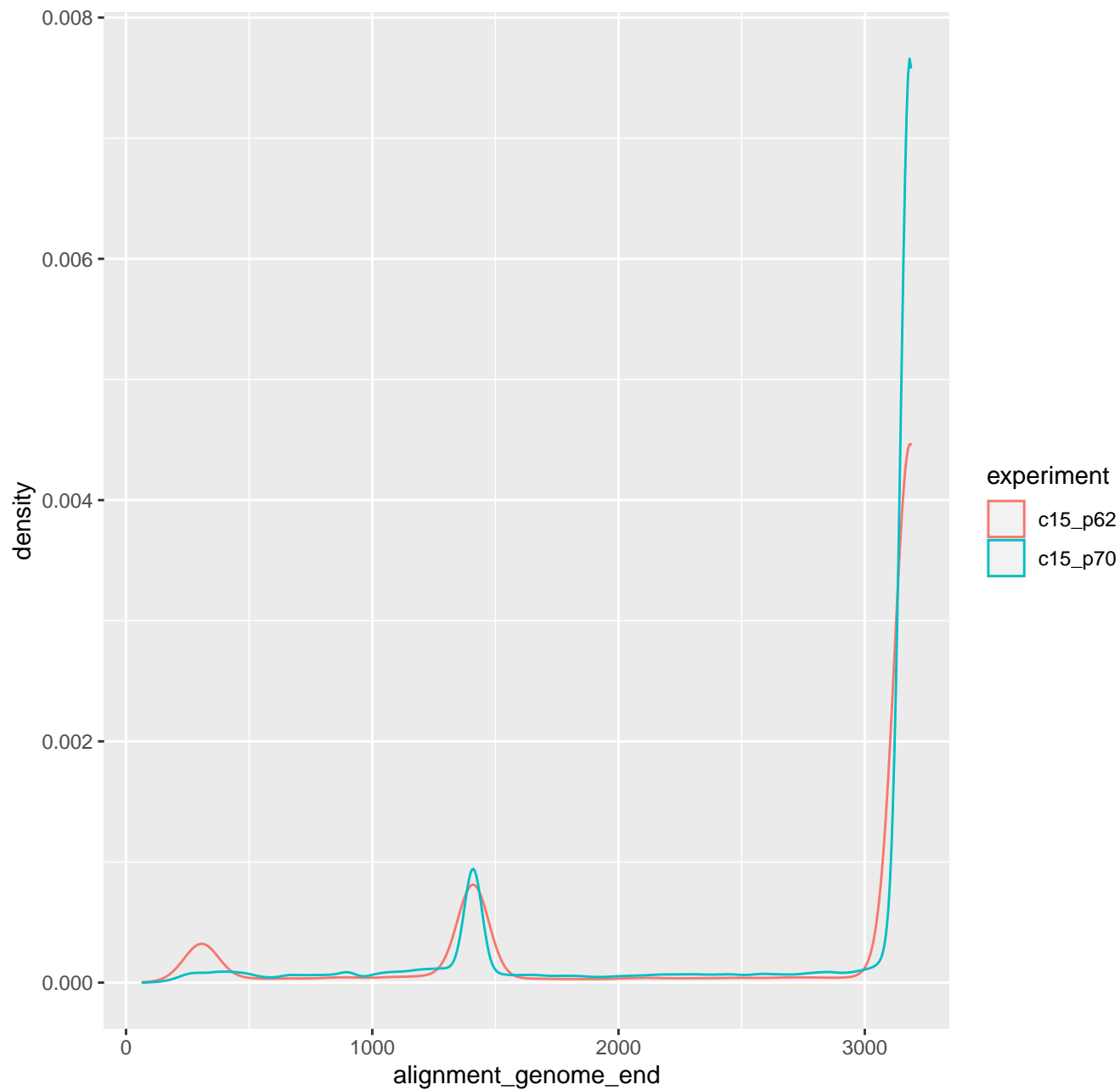
Alignment Coverage



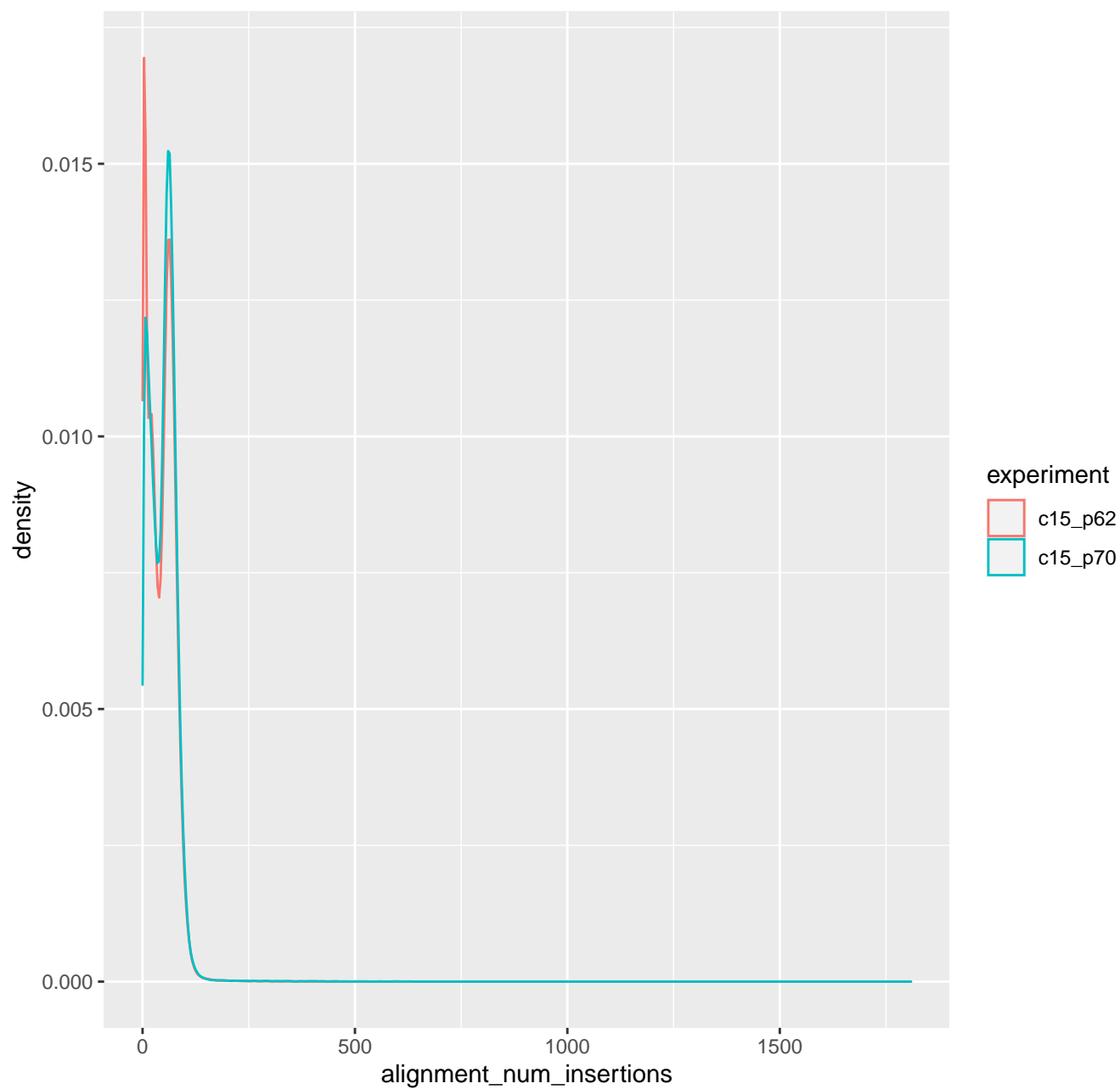
Alignment genome_start



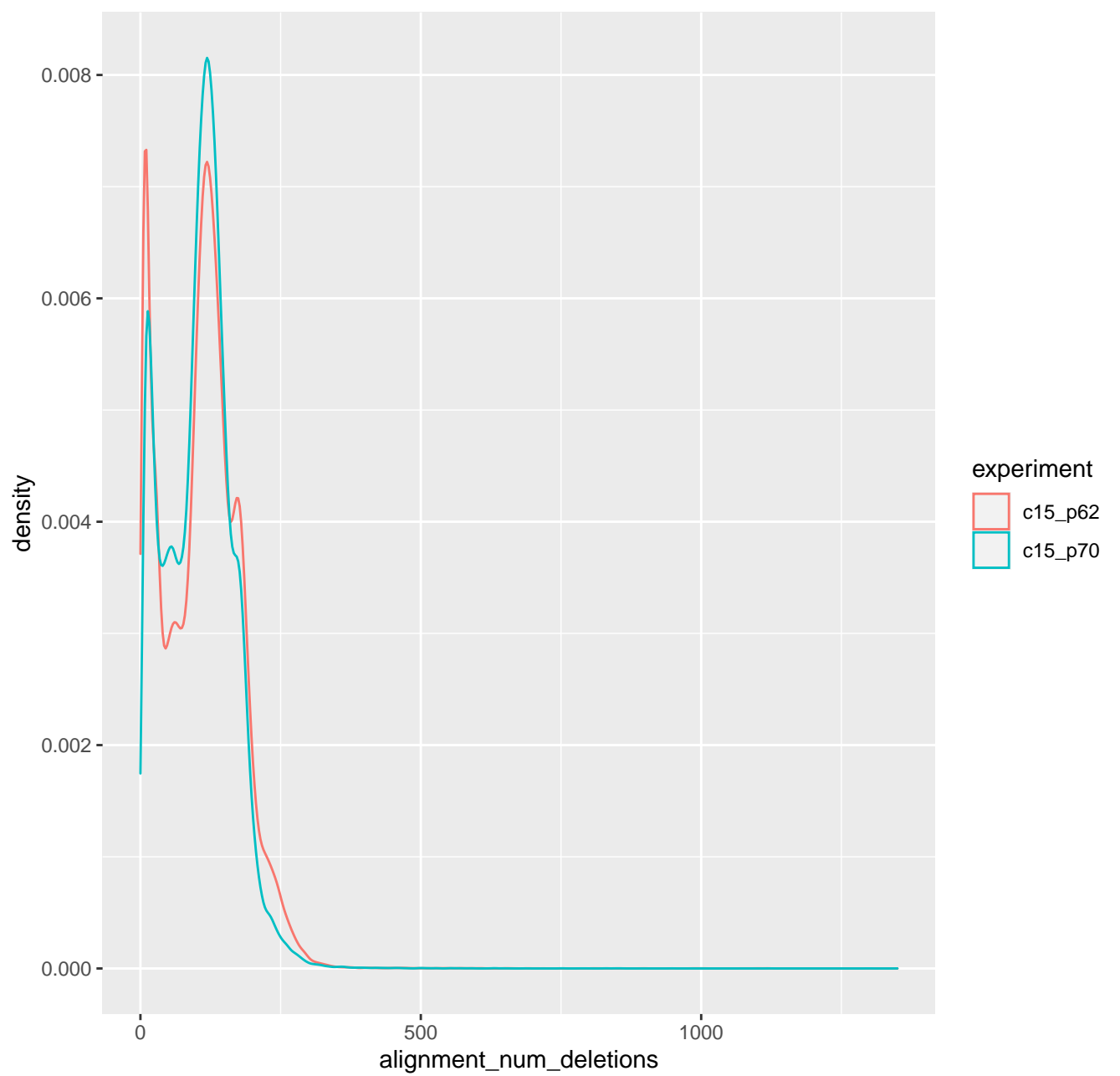
Alignment genome_end



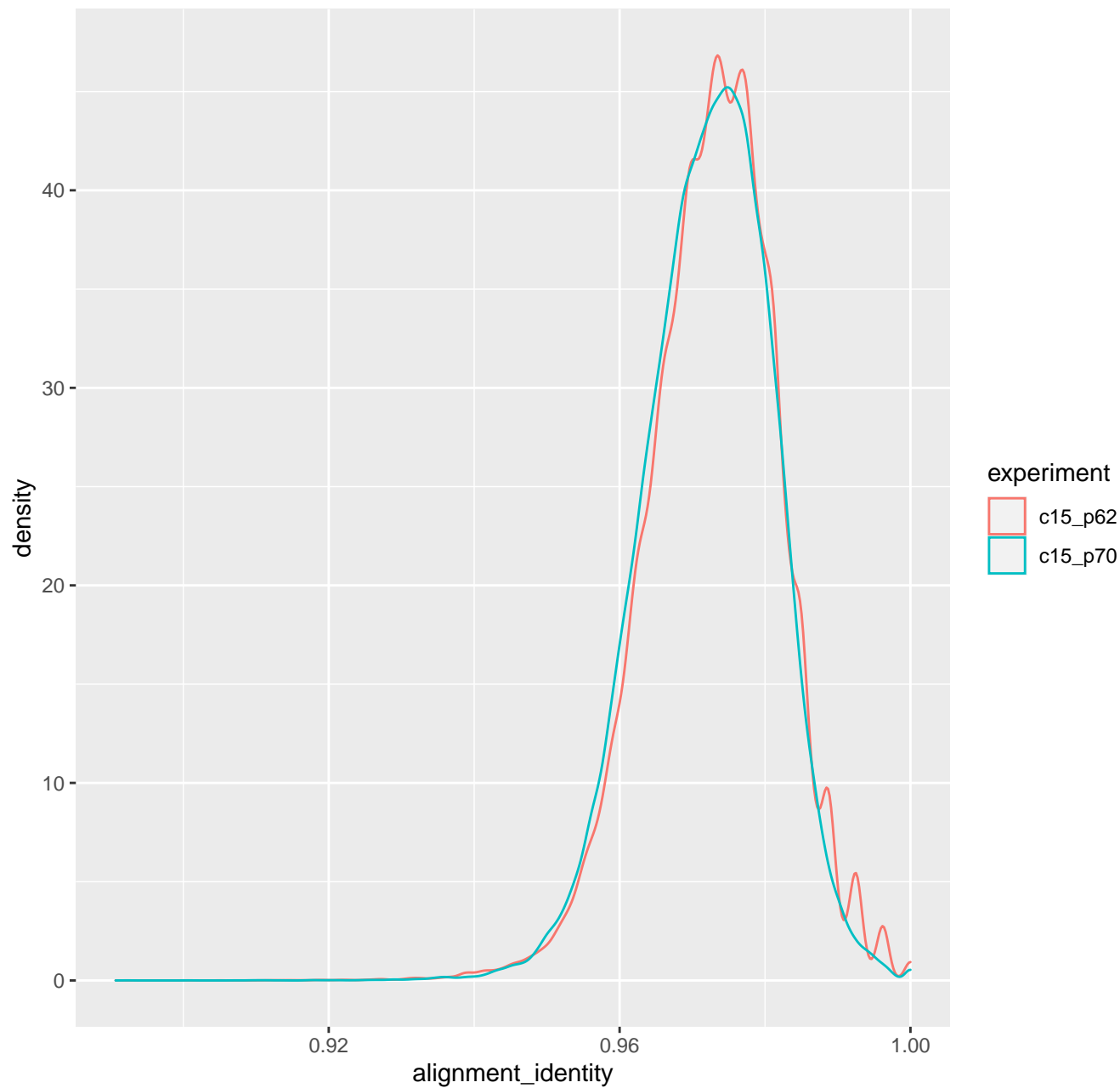
Number of insertions



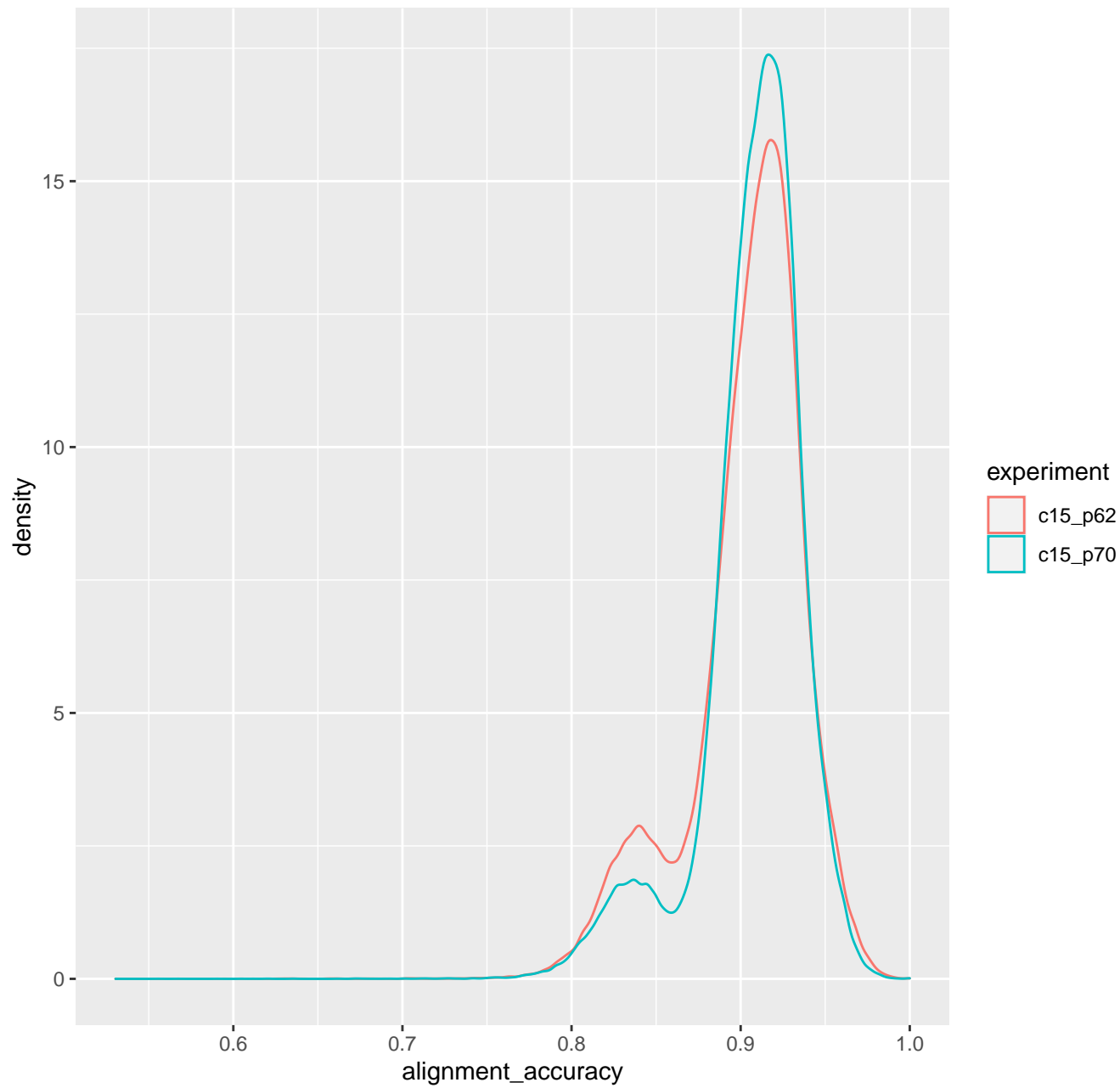
Number of deletions



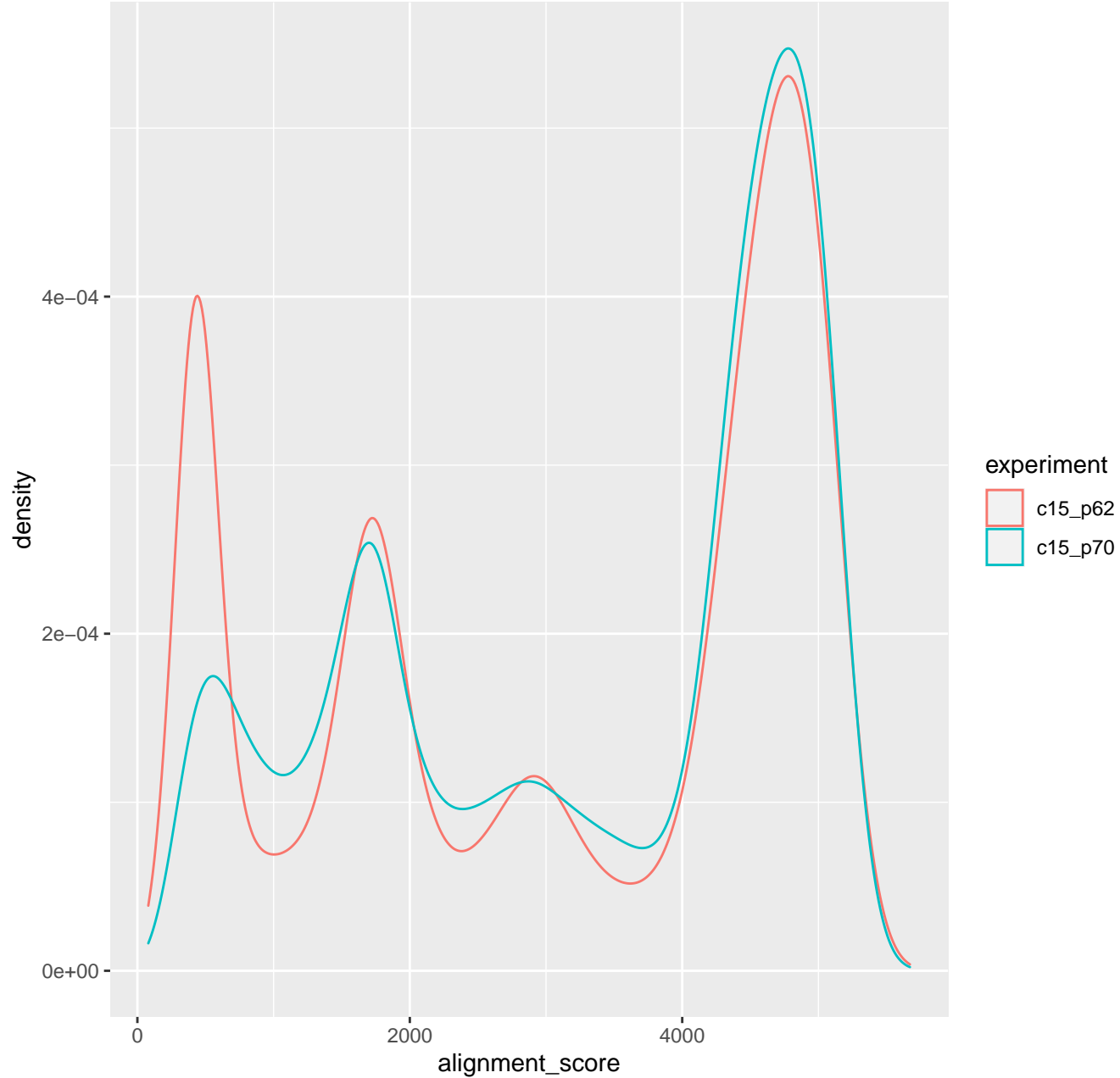
Alignment_identity



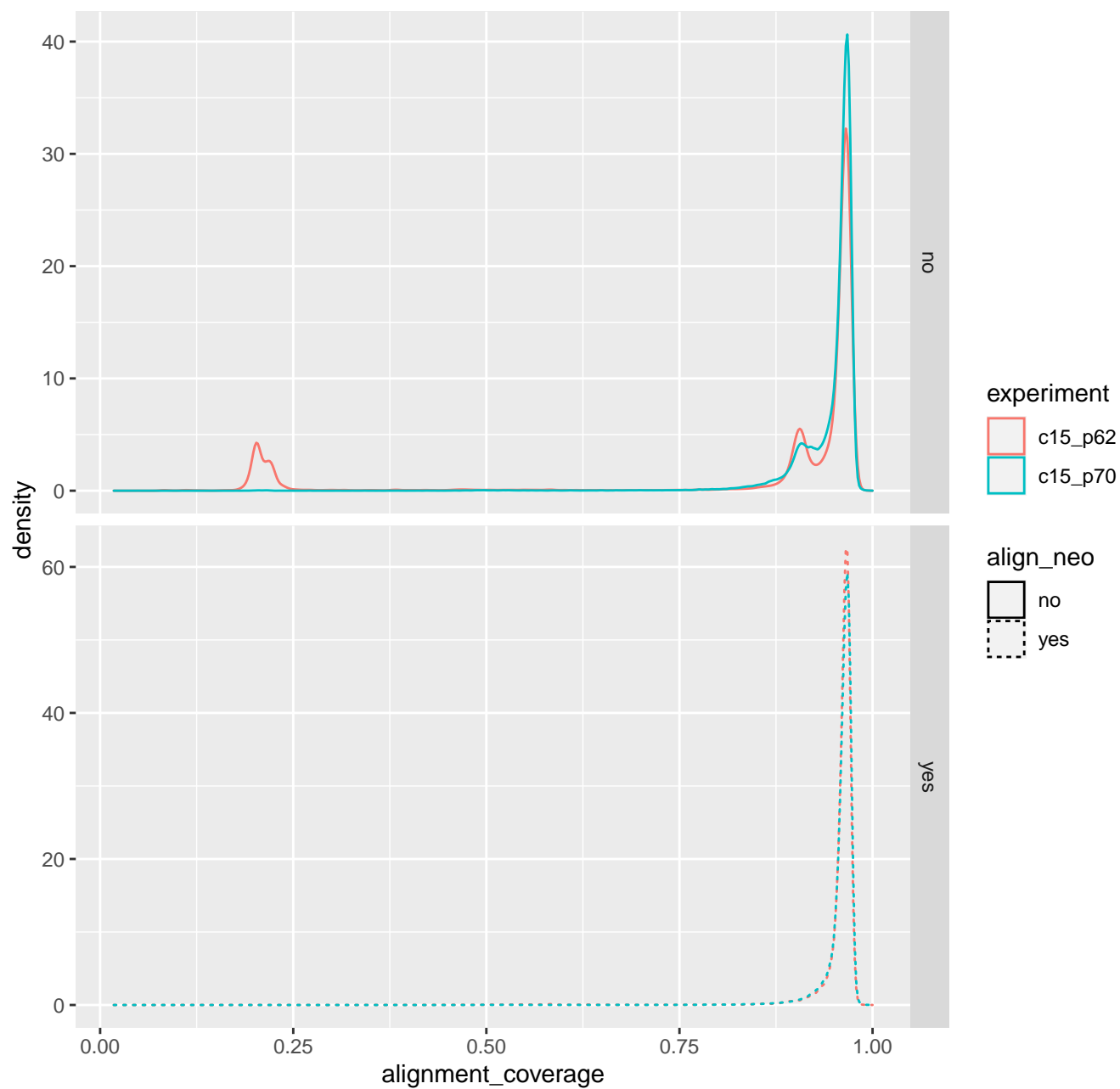
Alignment_accuracy



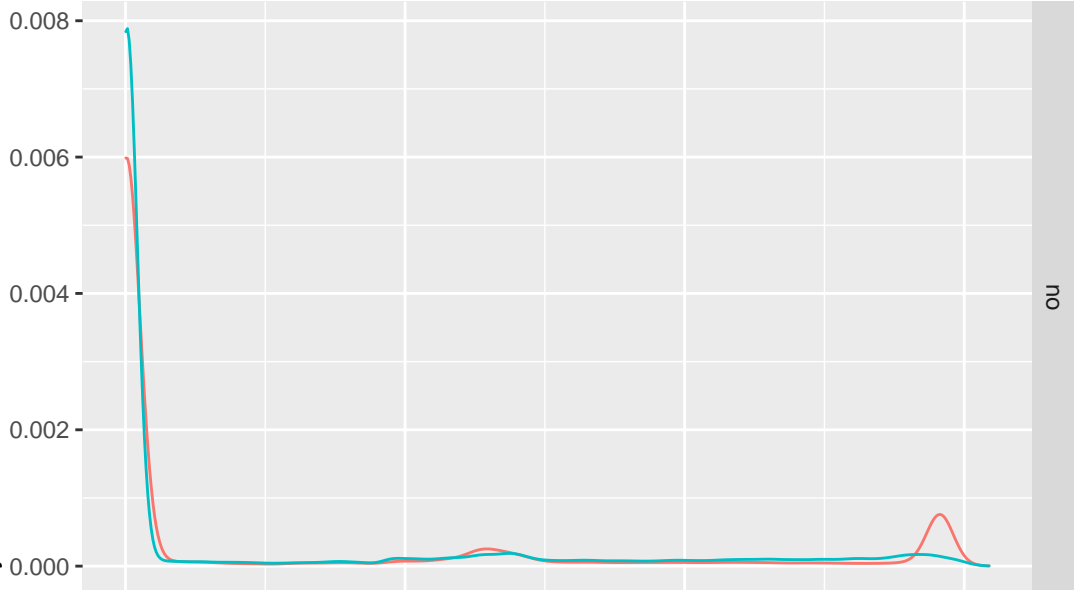
Alignment_score



Alignment Coverage



Alignment genome_start



experiment

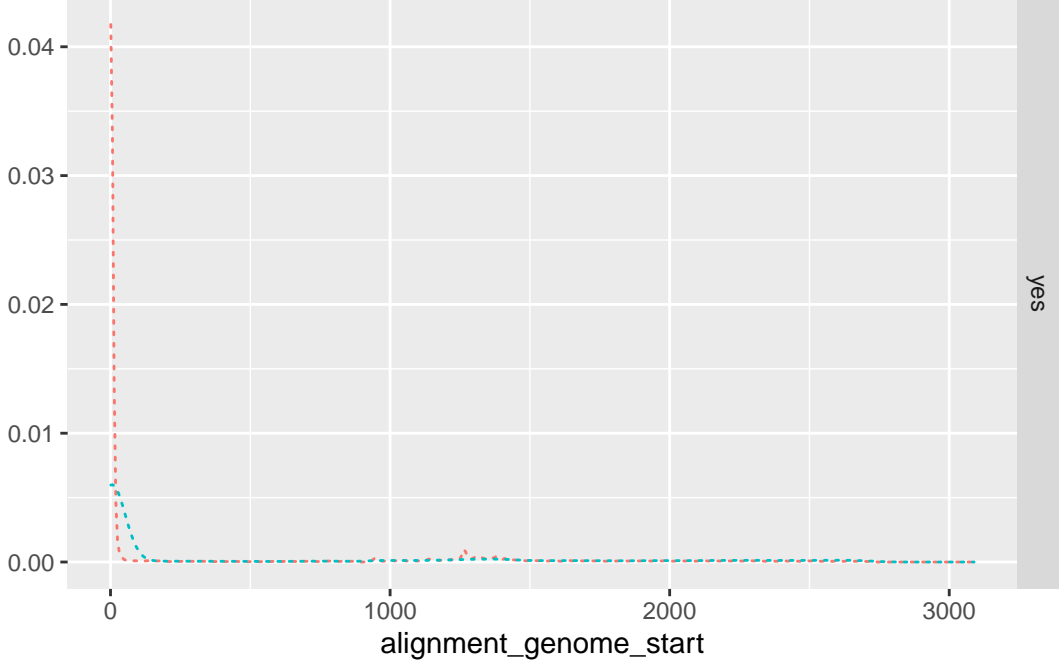
c15_p62

c15_p70

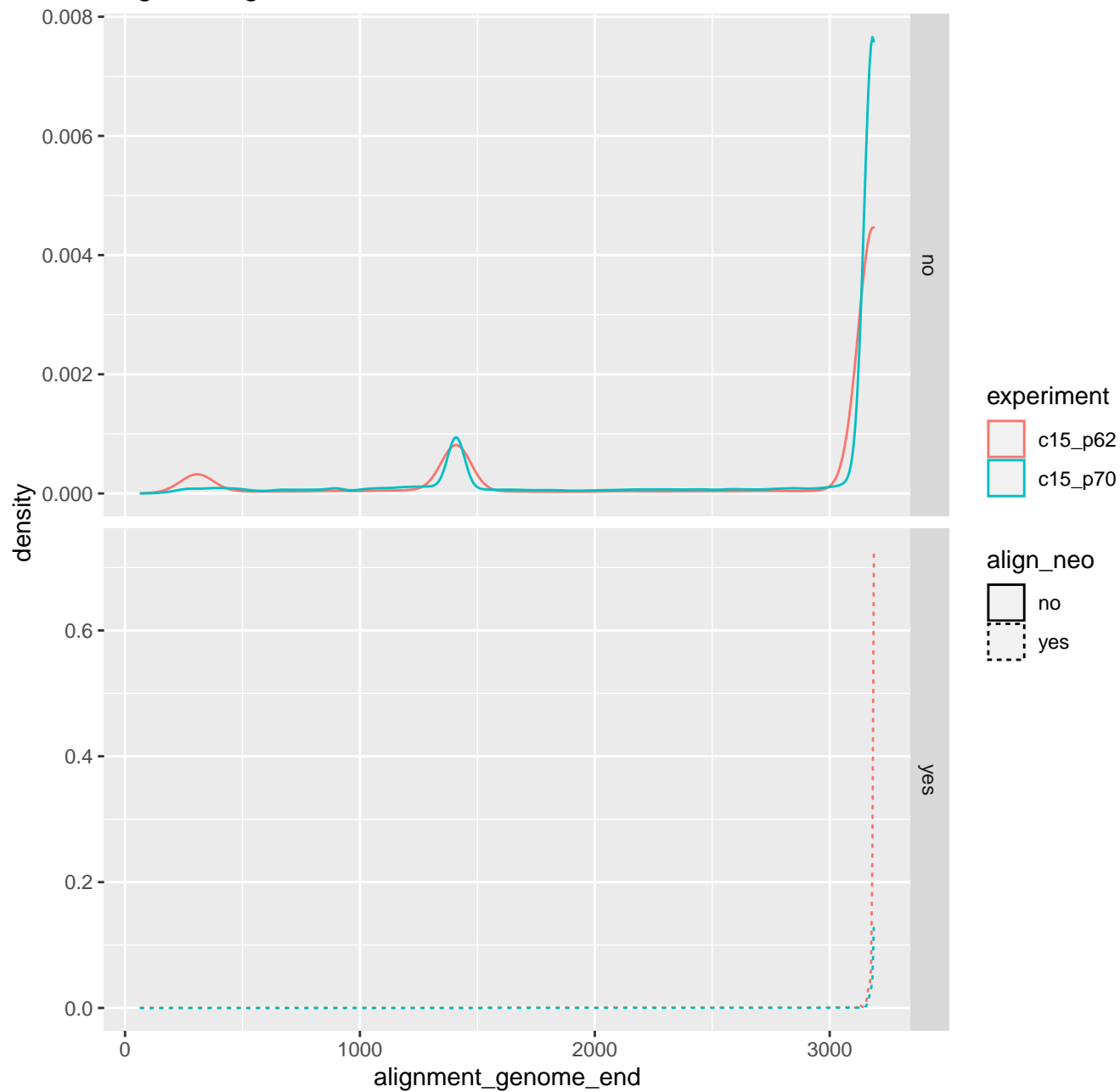
align_neo

no

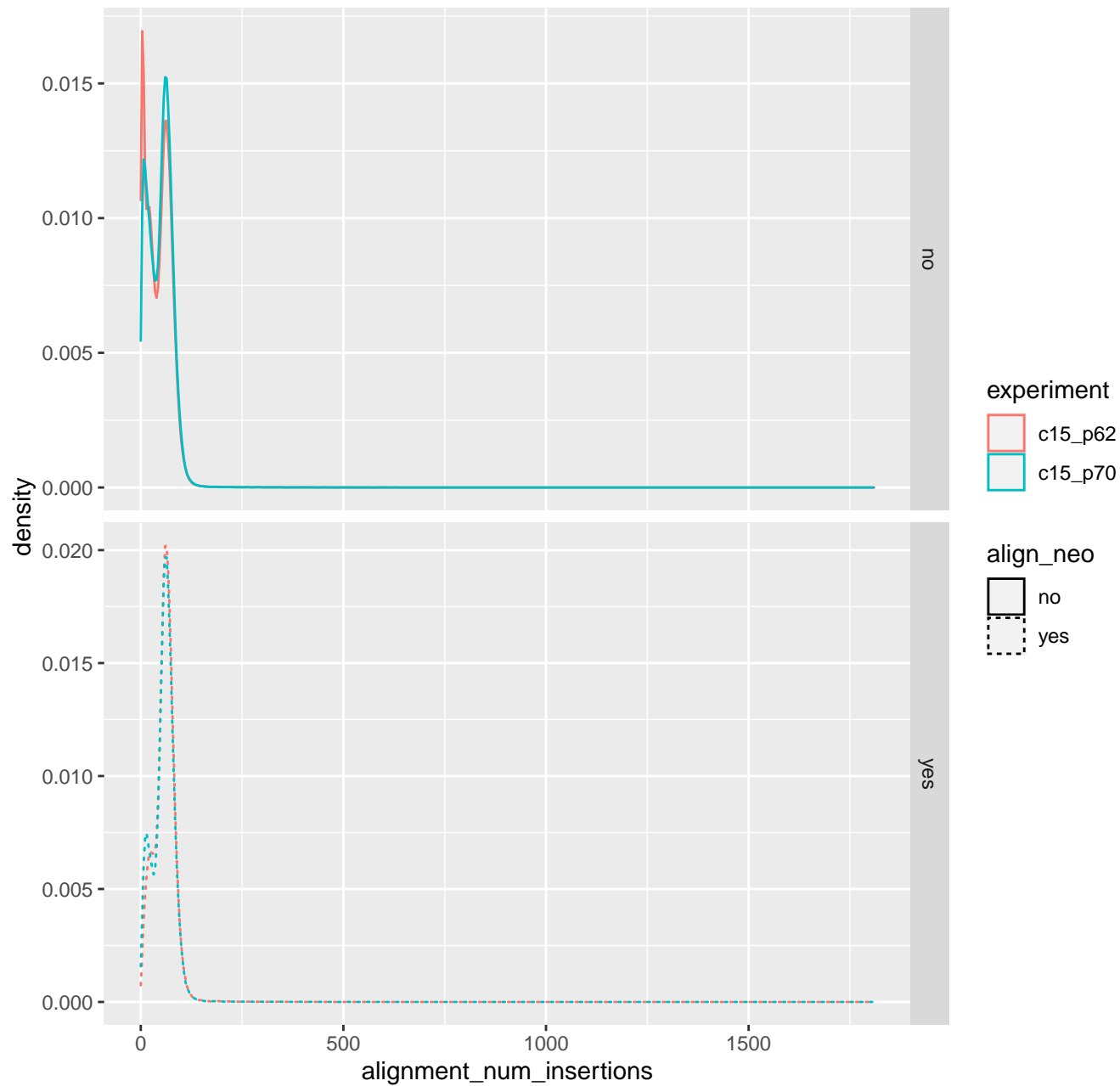
yes



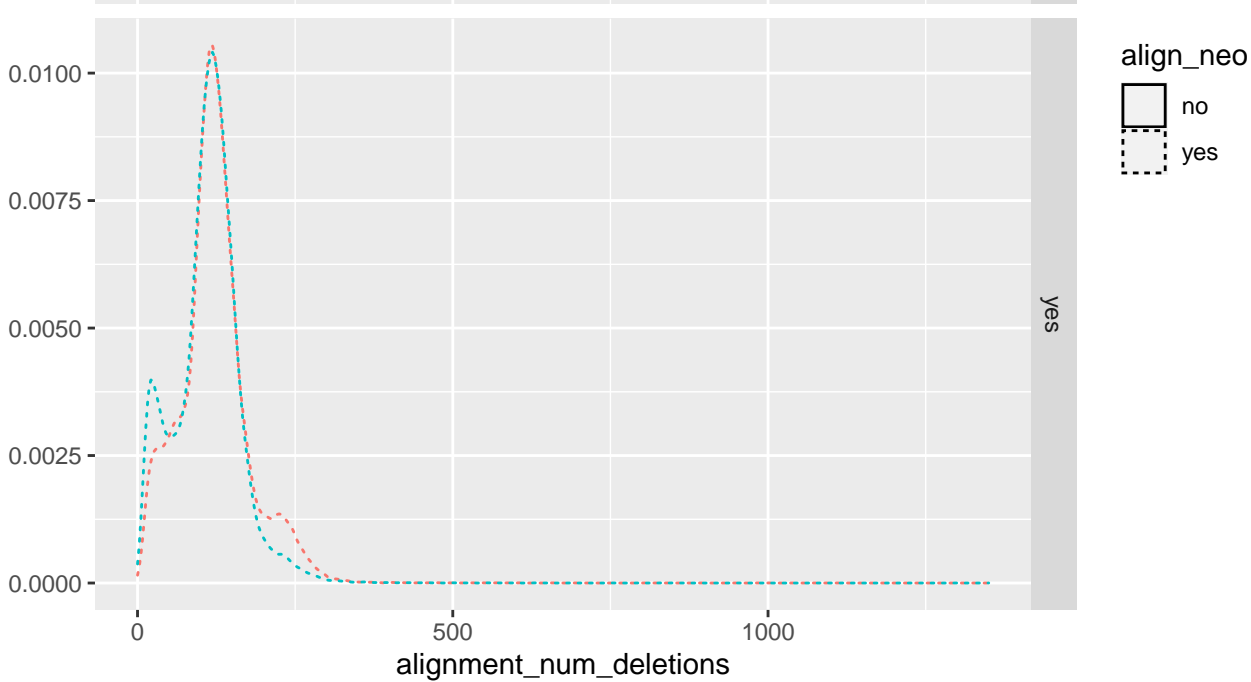
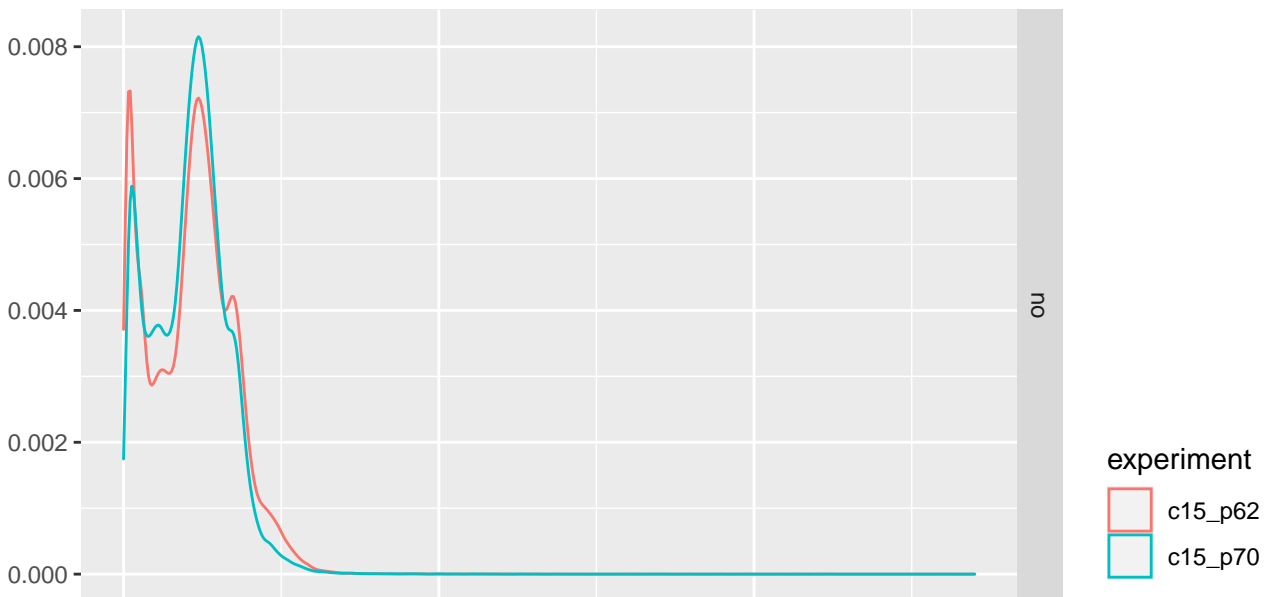
Alignment genome_end



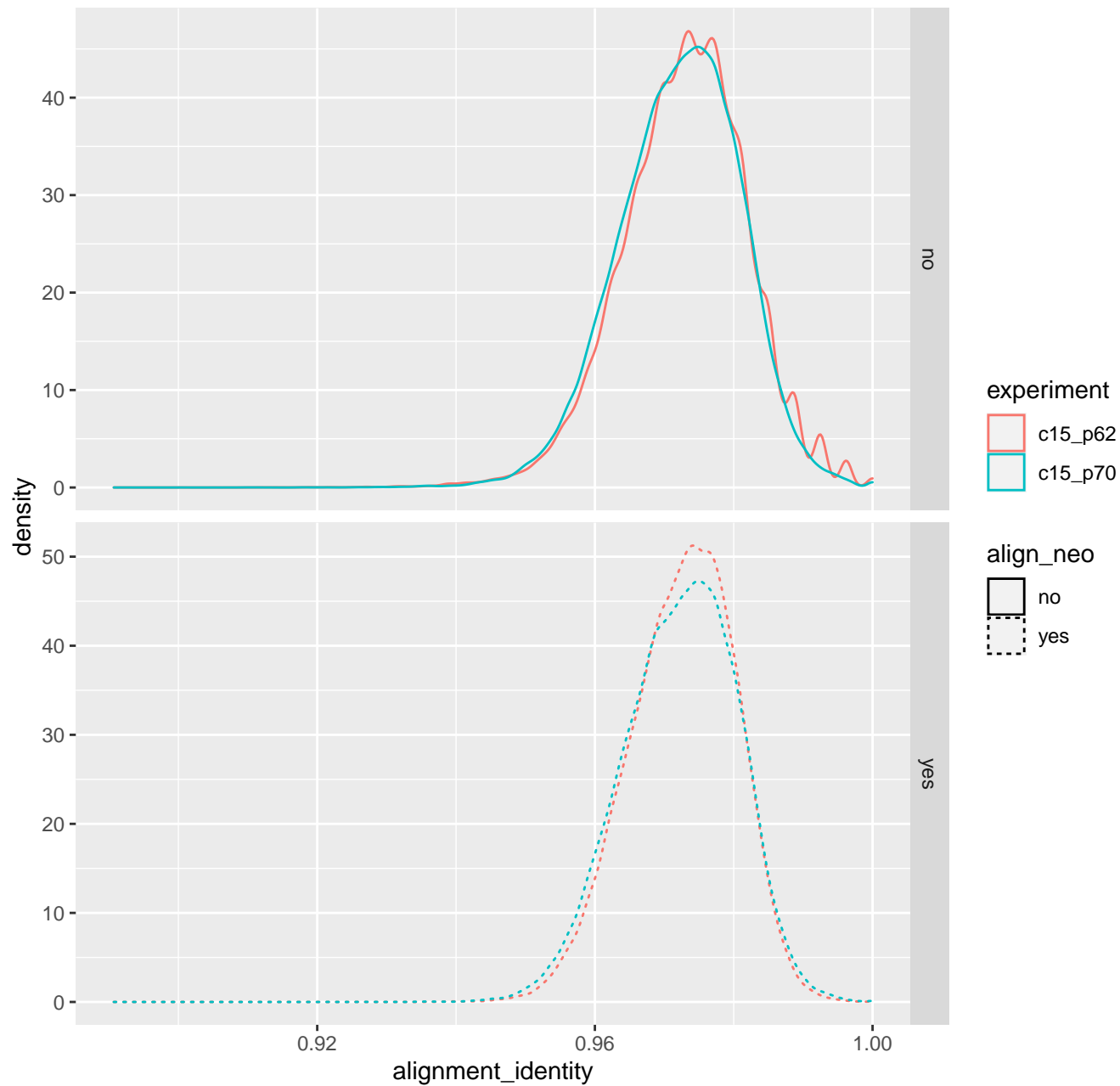
Number of insertions



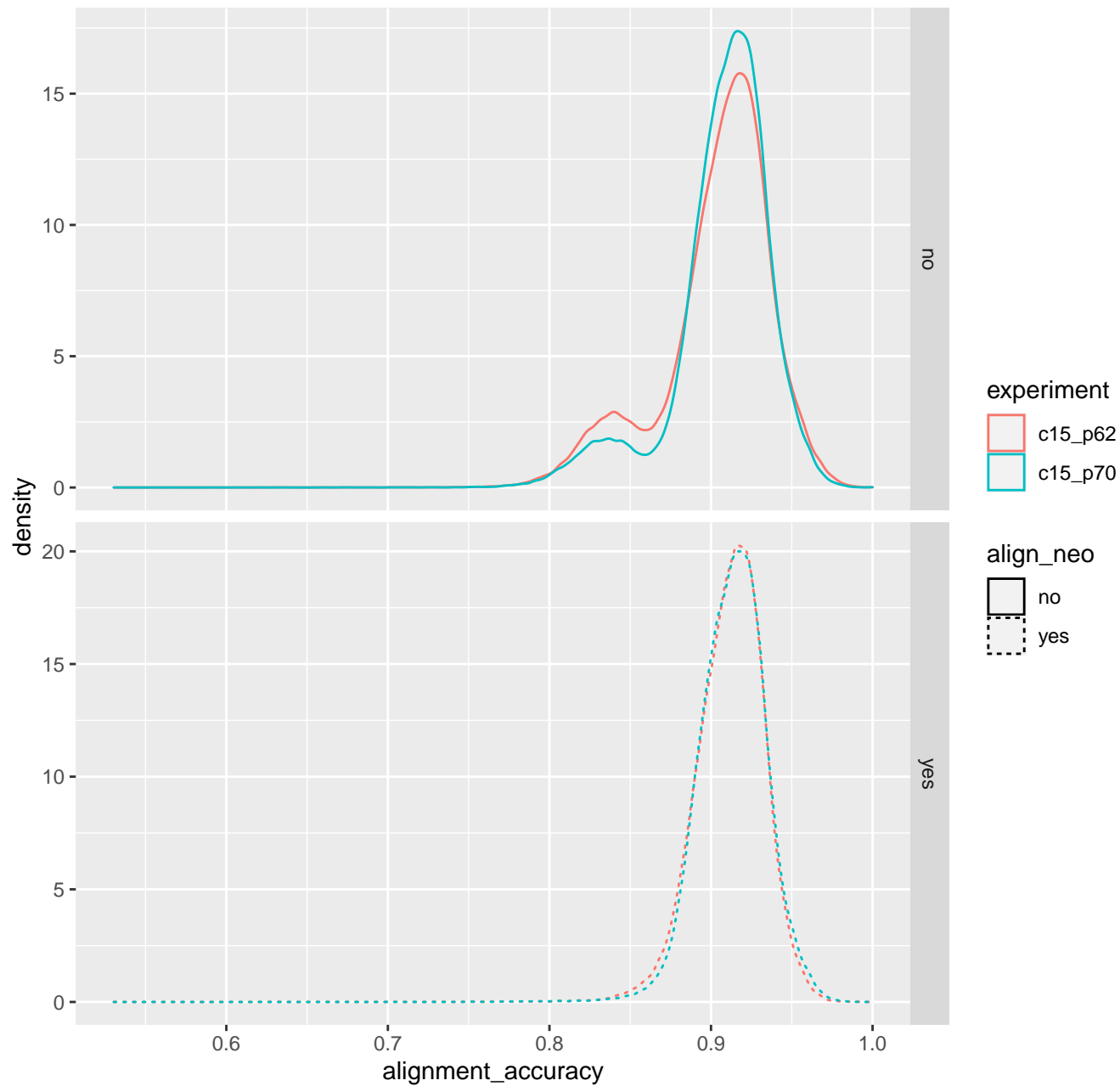
Number of deletions



Alignment_identity



Alignment_accuracy



Alignment_score

