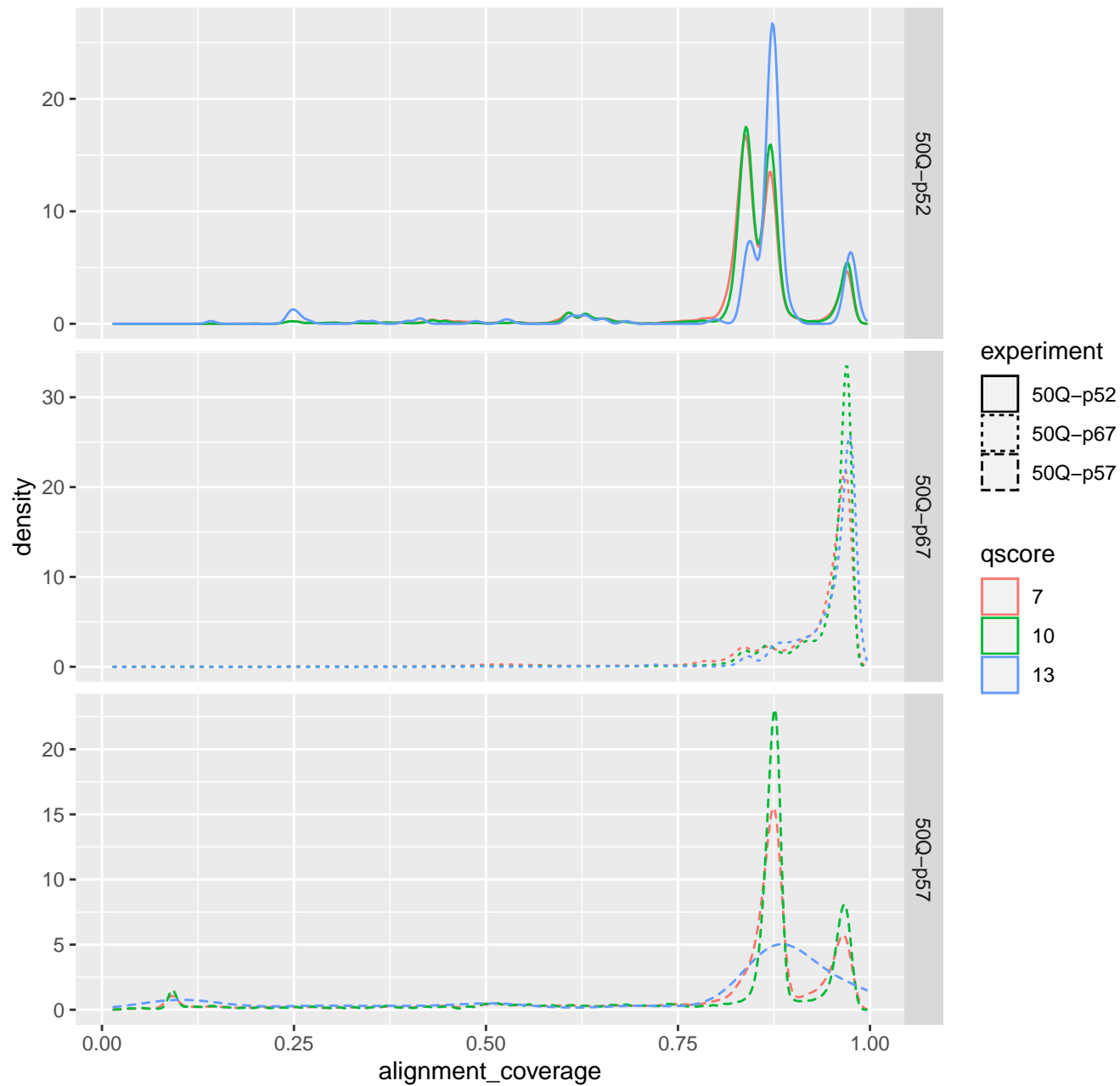
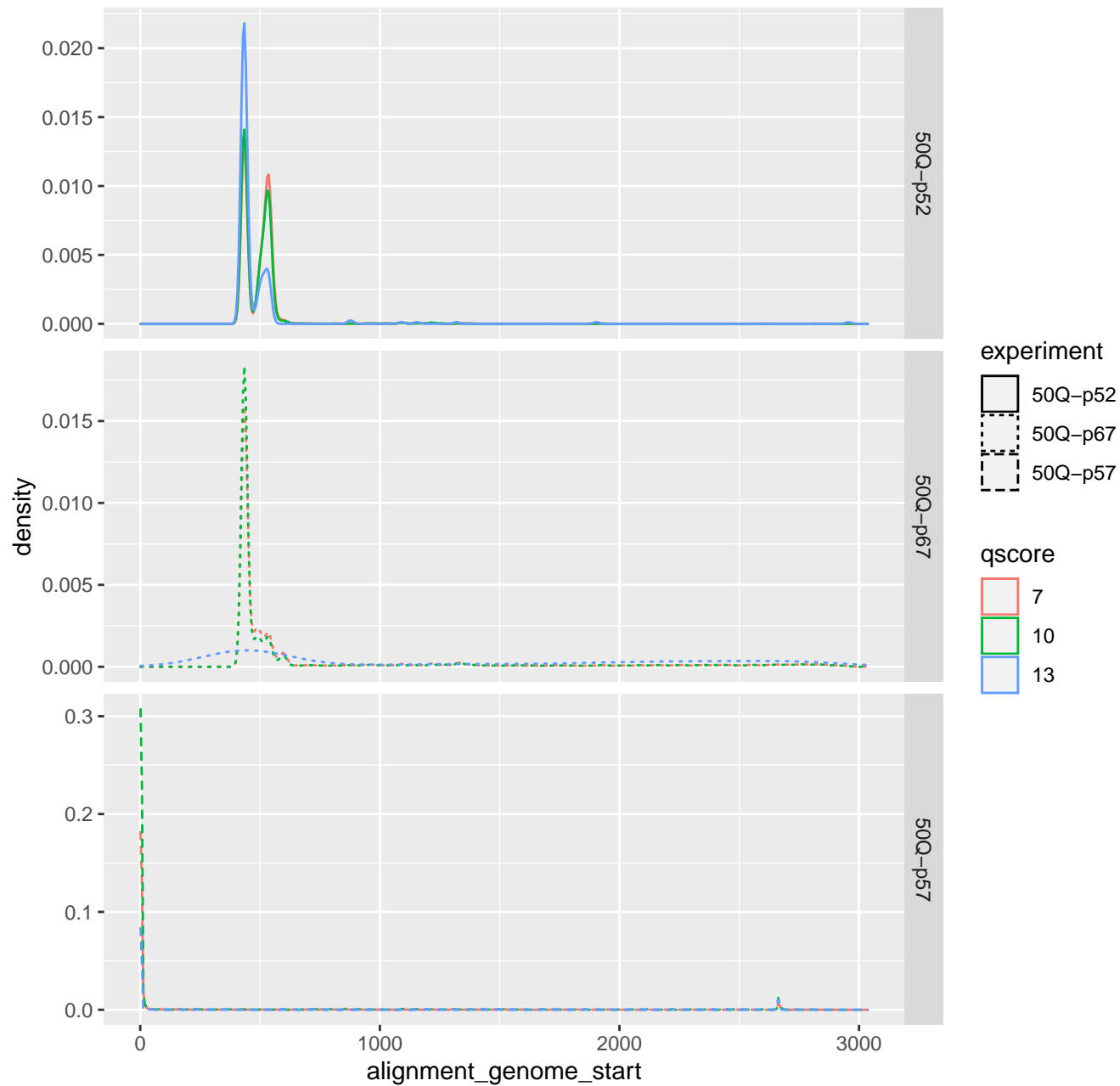


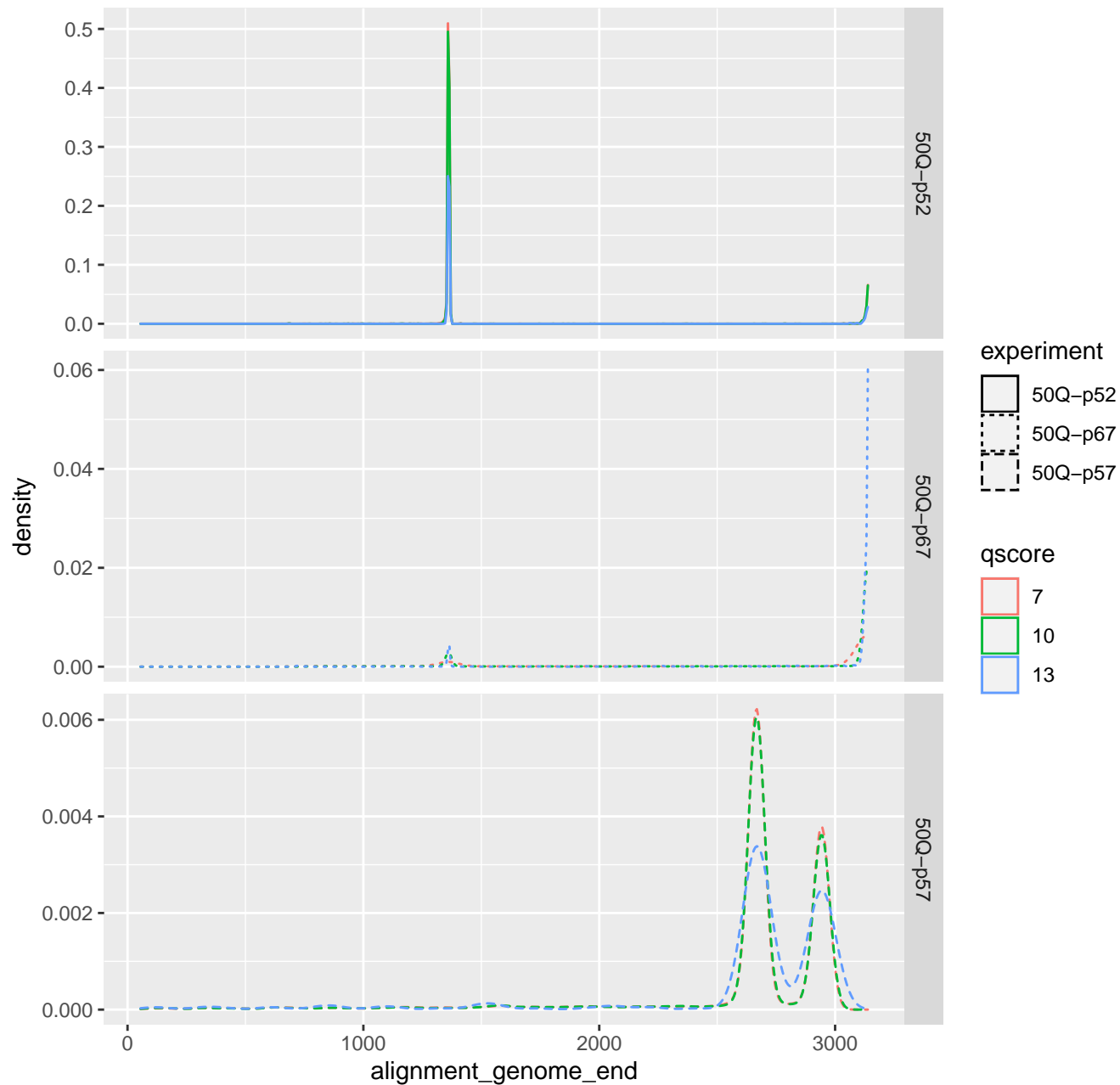
Alignment Coverage for each Qscore value



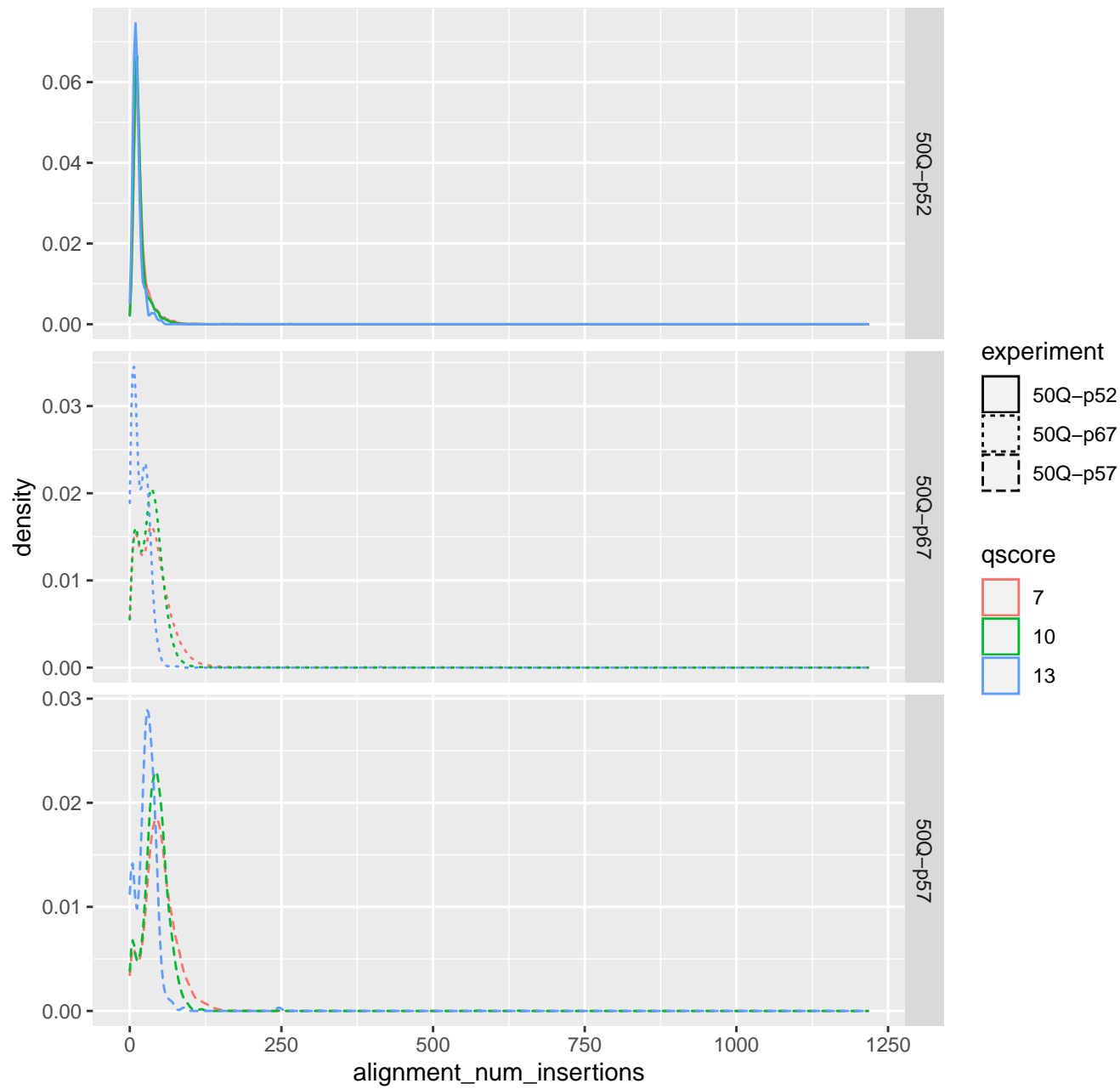
Alignment genome_start for each Qscore value



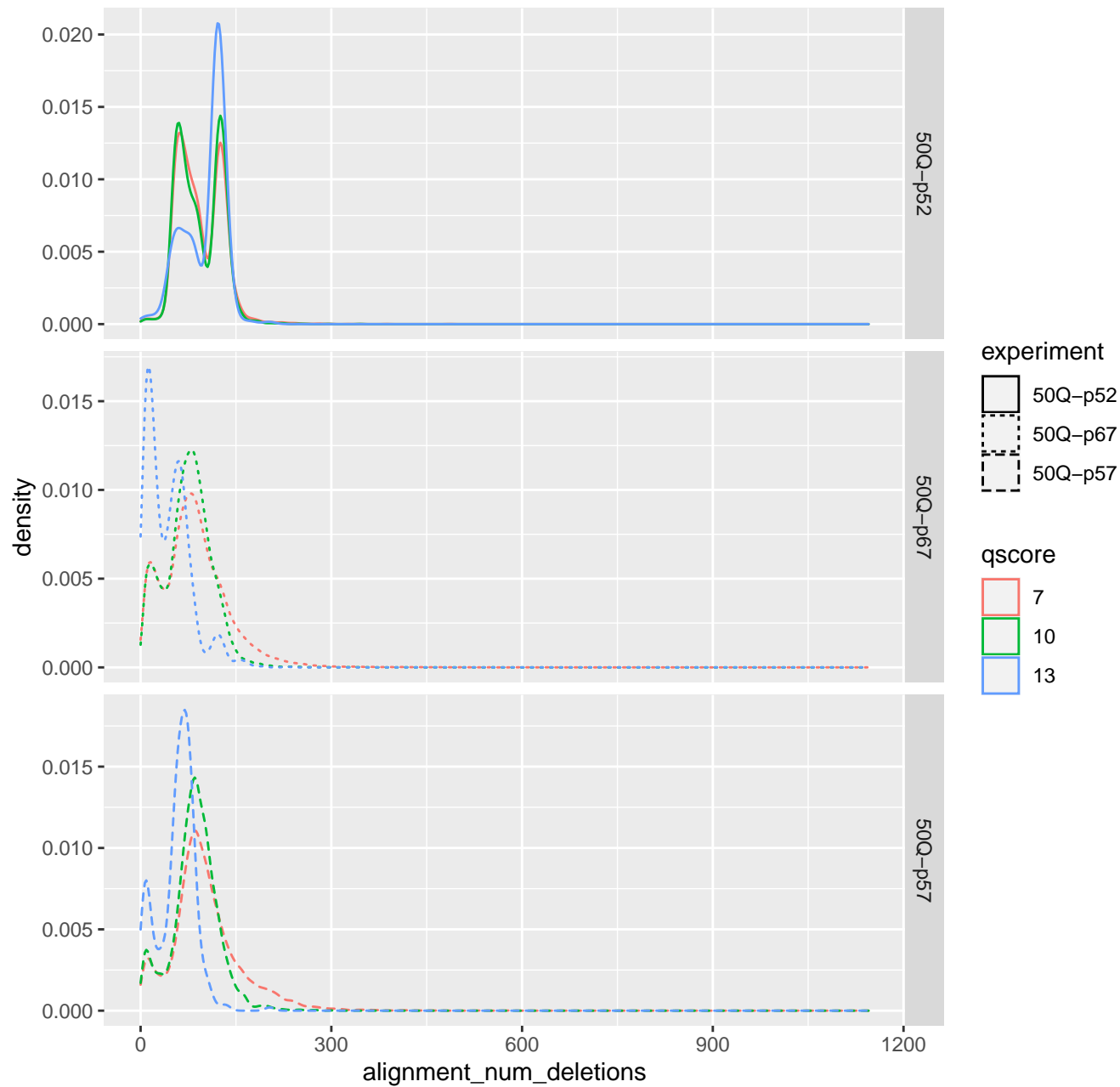
Alignment genome_end for each Qscore value



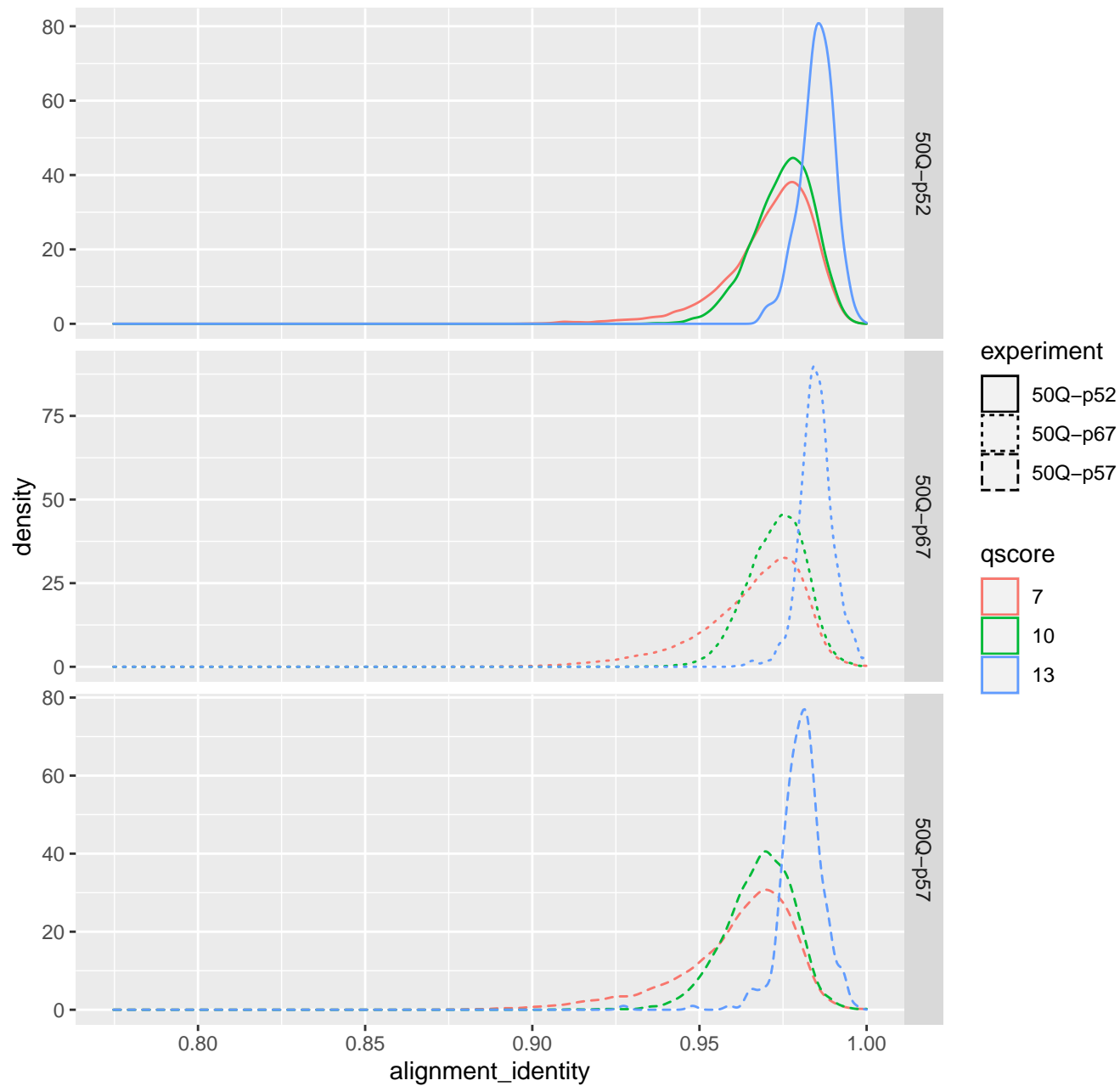
Number of insertions for each Qscore value



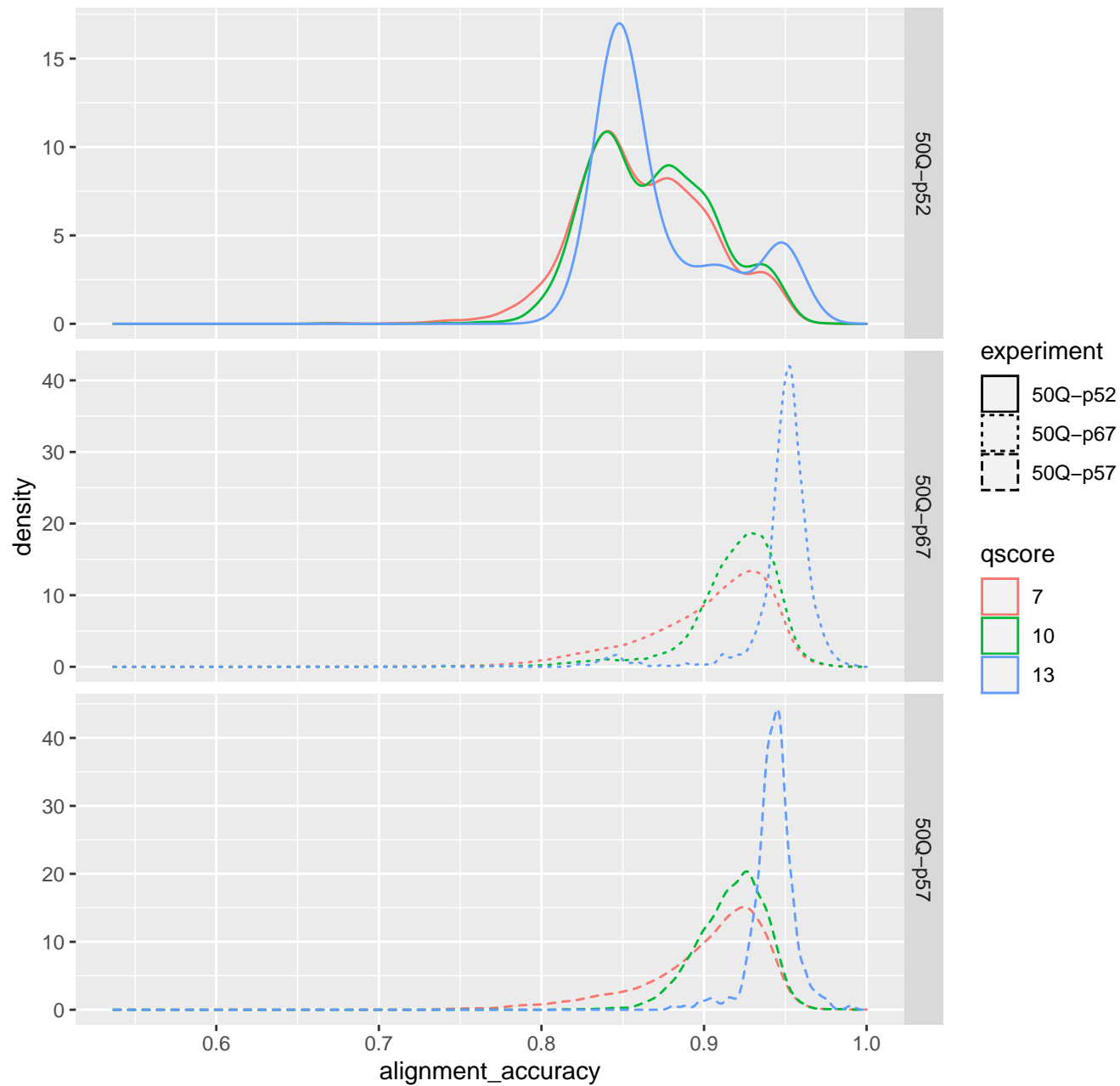
Number of deletions for each Qscore value



Alignment_identity for each Qscore value



Alignment_accuracy for each Qscore value



Alignment_score for each Qscore value

