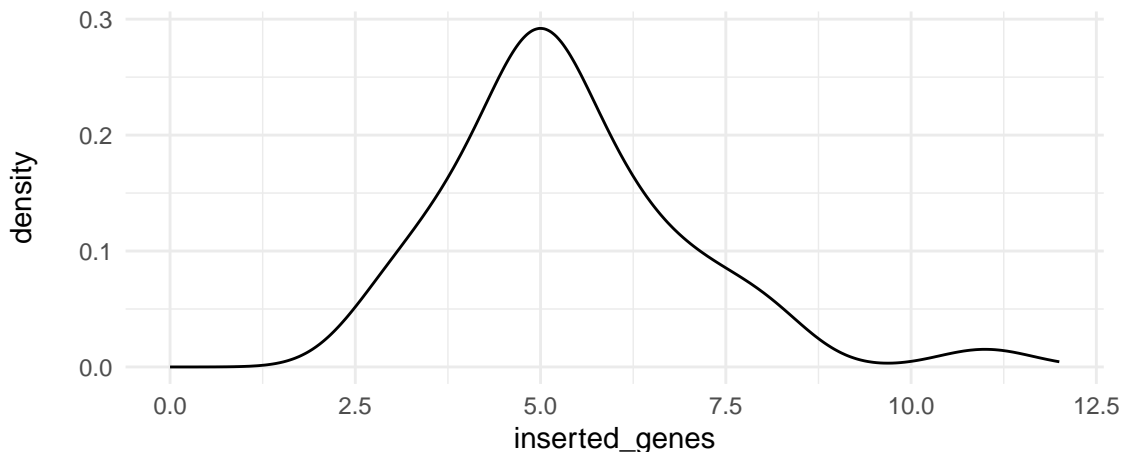


# number of inserted genes between ORF10 and ORF17

complete relatives only (n = 41)



# length (bp) of inserted region between ORF10 and ORF17

complete relatives only (n = 41)

