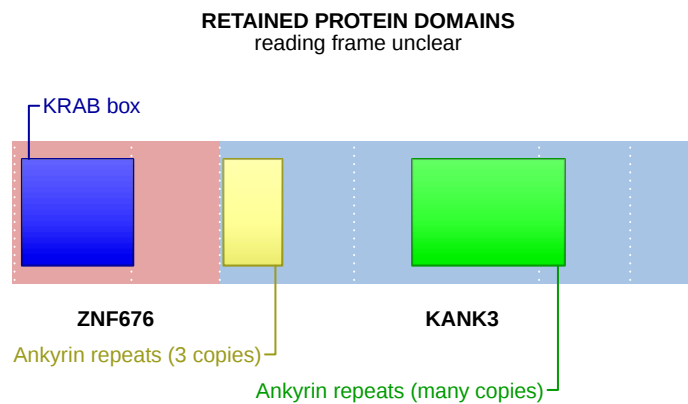
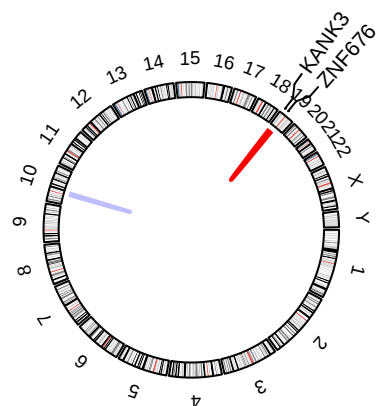
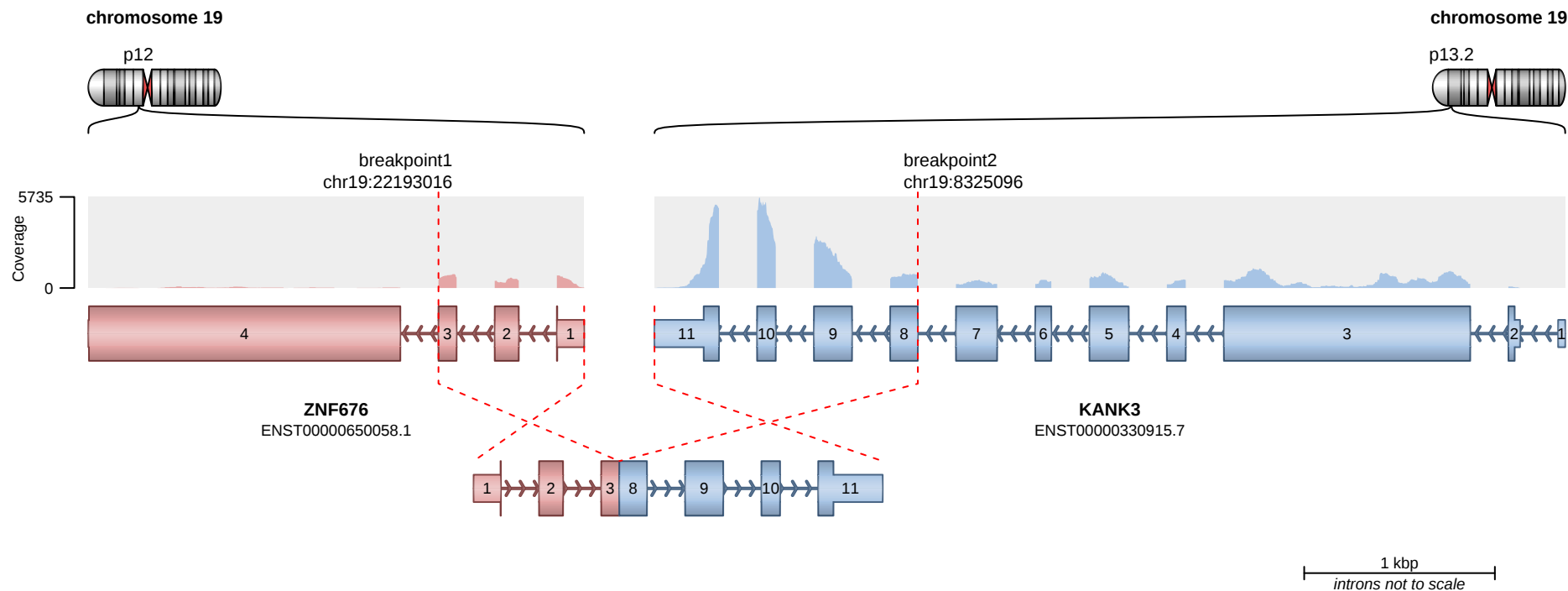
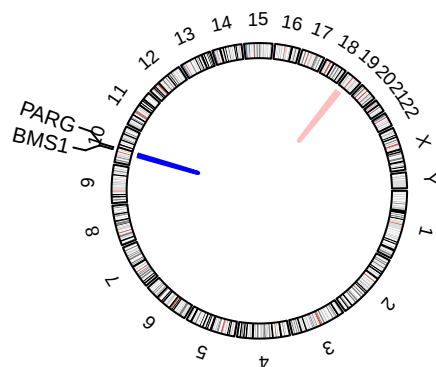
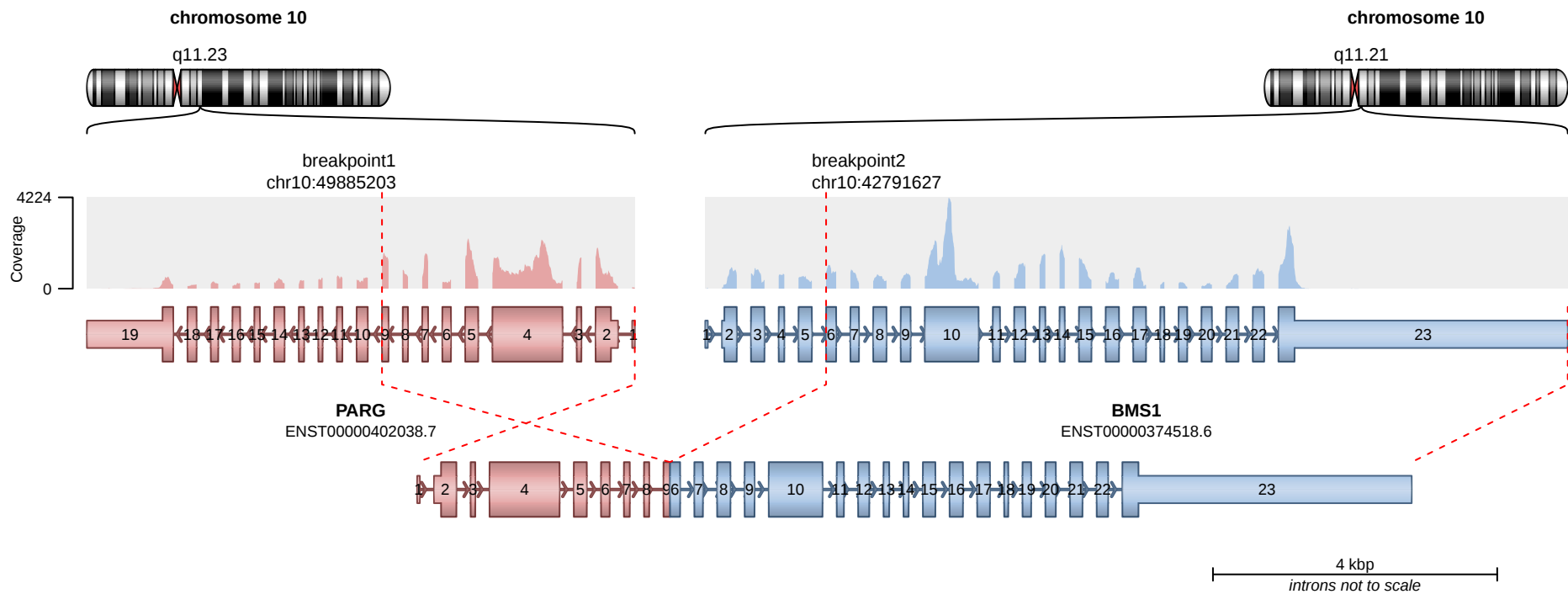




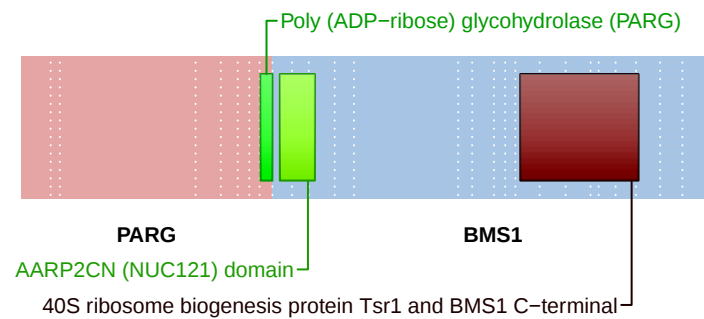
SUPPORTING READ COUNT

Split reads = 116
Discordant mates = 0

translocation
duplication
deletion
inversion



RETAINED PROTEIN DOMAINS reading frame unclear



SUPPORTING READ COUNT

Split reads = 62
Discordant mates = 0

— translocation — deletion
— duplication — inversion