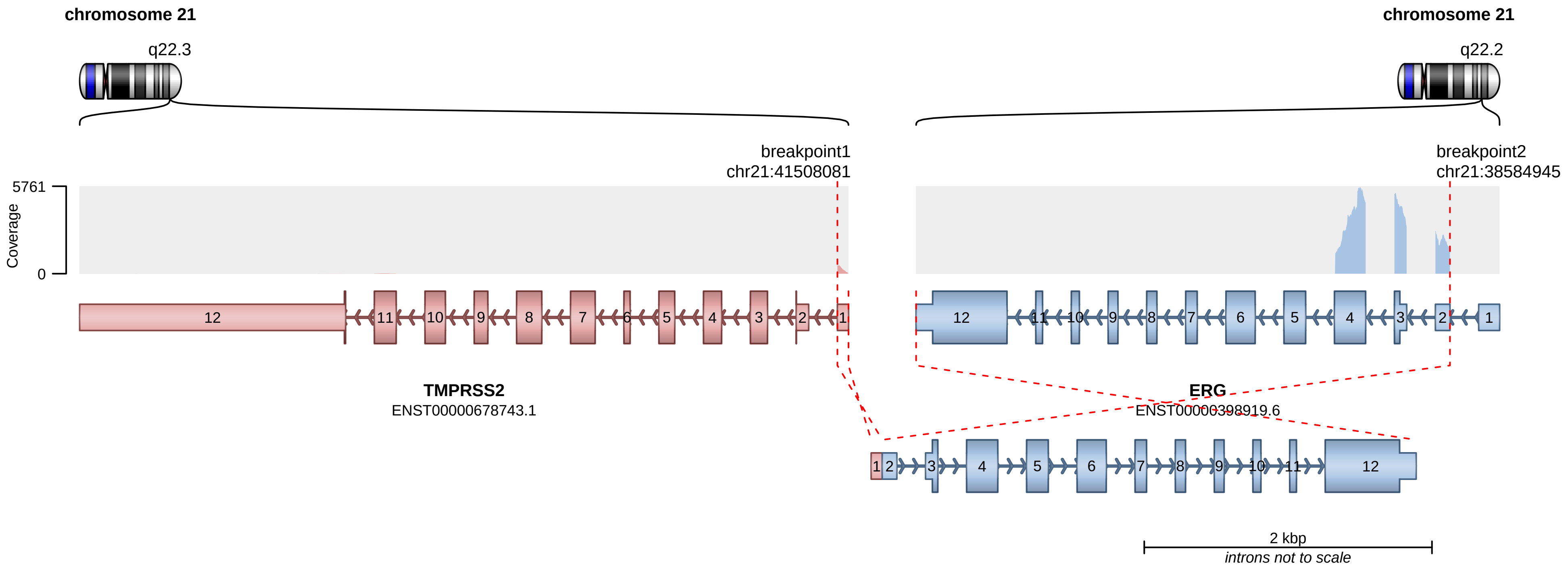
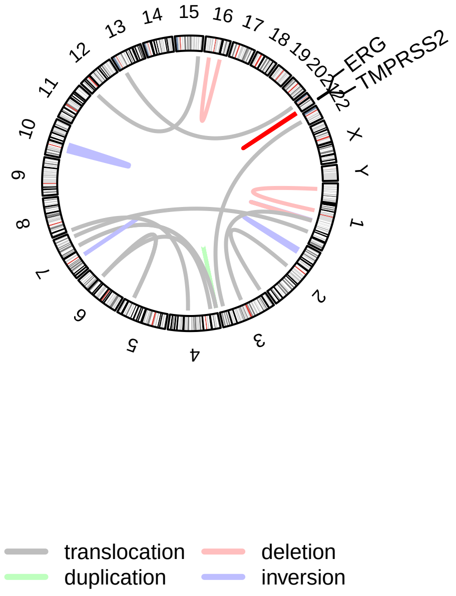
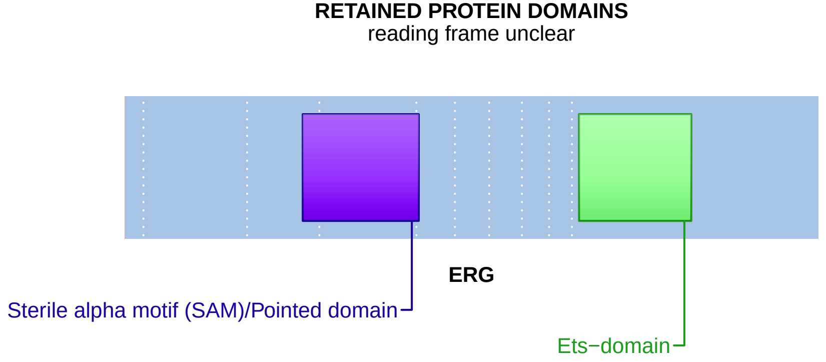
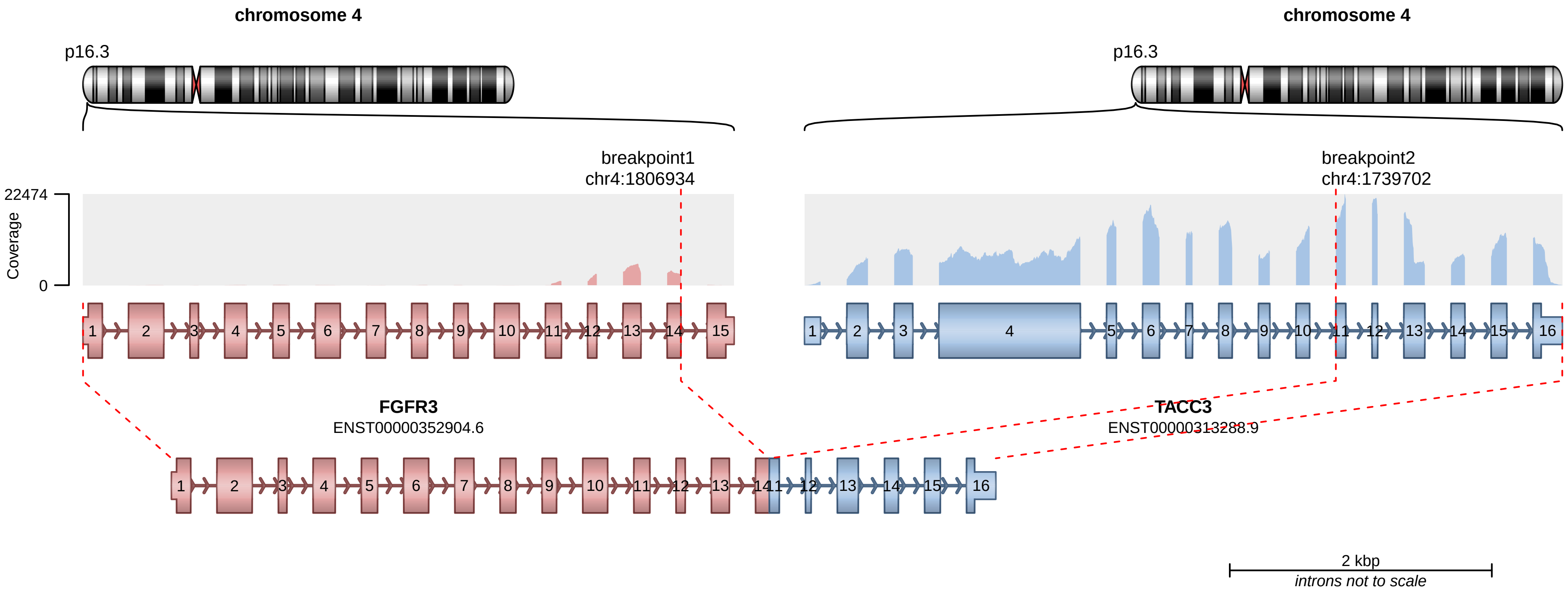


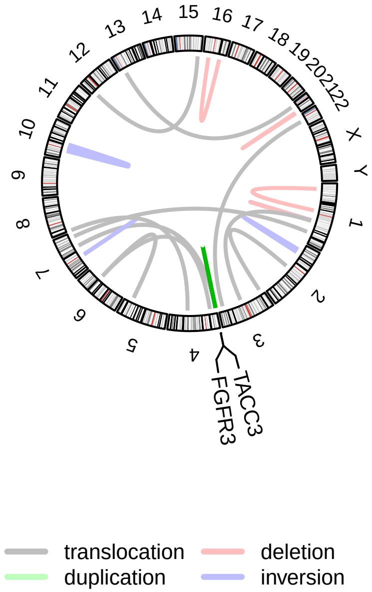


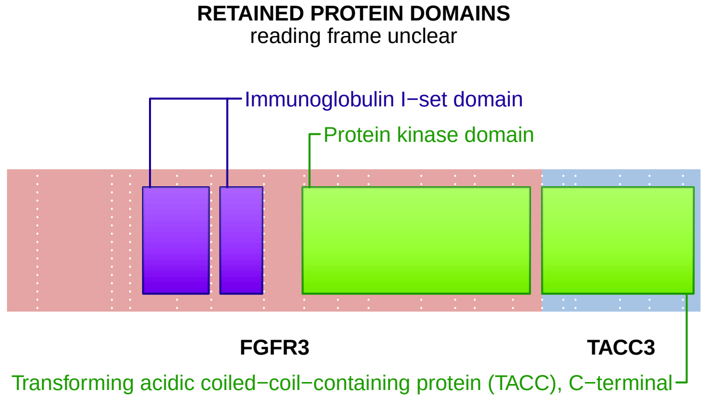


SUPPORTING READ COUNT

Split reads = 618
Discordant mates = 0

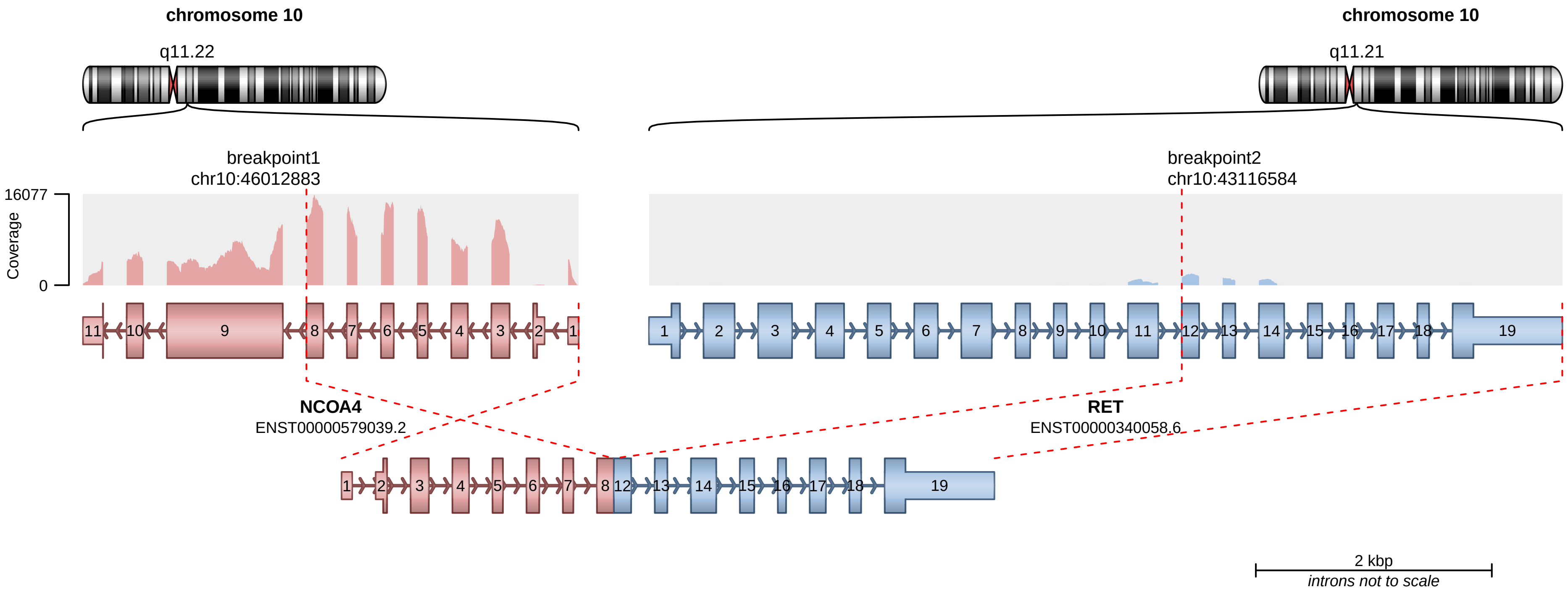


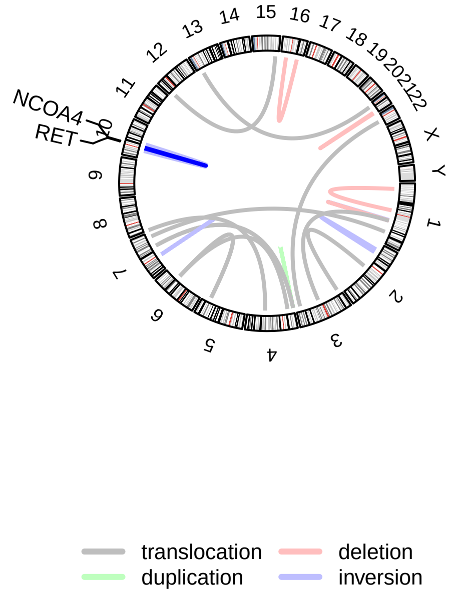




SUPPORTING READ COUNT

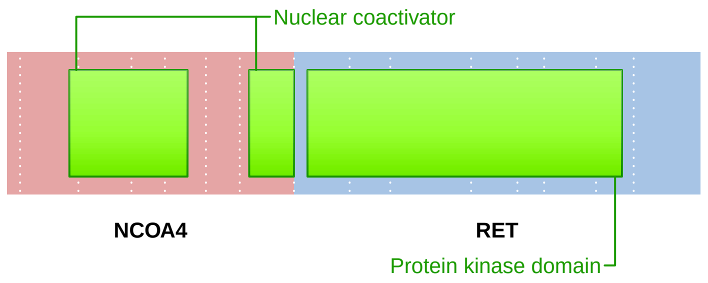
Split reads = 551
Discordant mates = 0





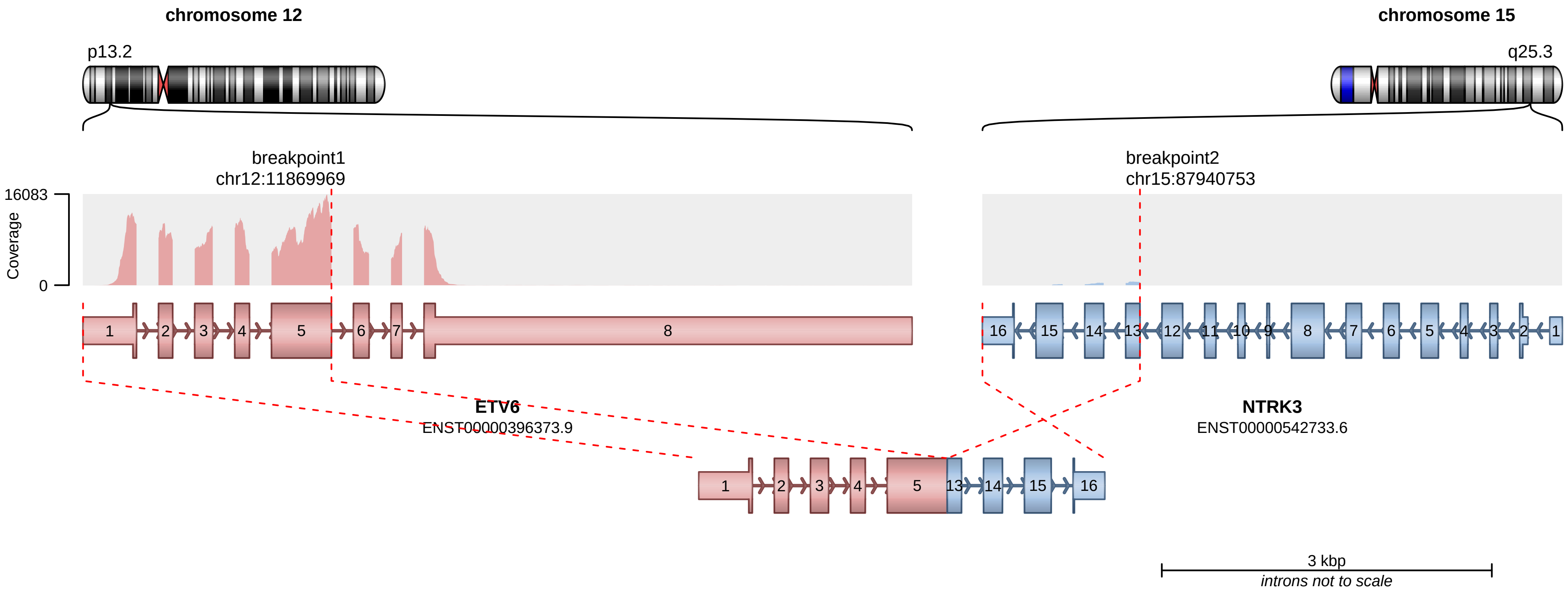
RETAINED PROTEIN DOMAINS

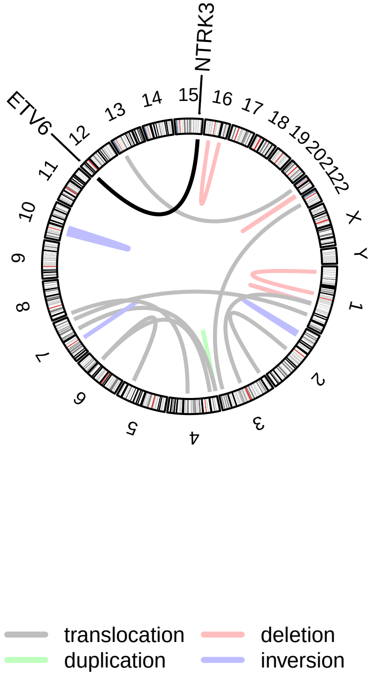
reading frame unclear



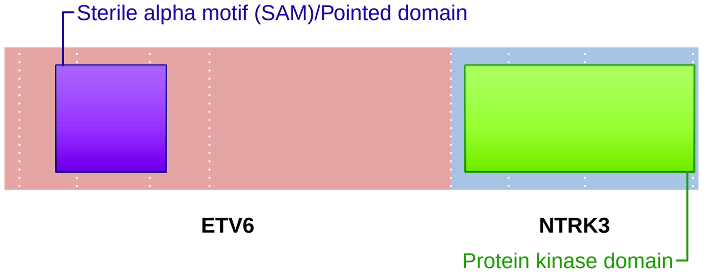
SUPPORTING READ COUNT

Split reads = 219
Discordant mates = 0



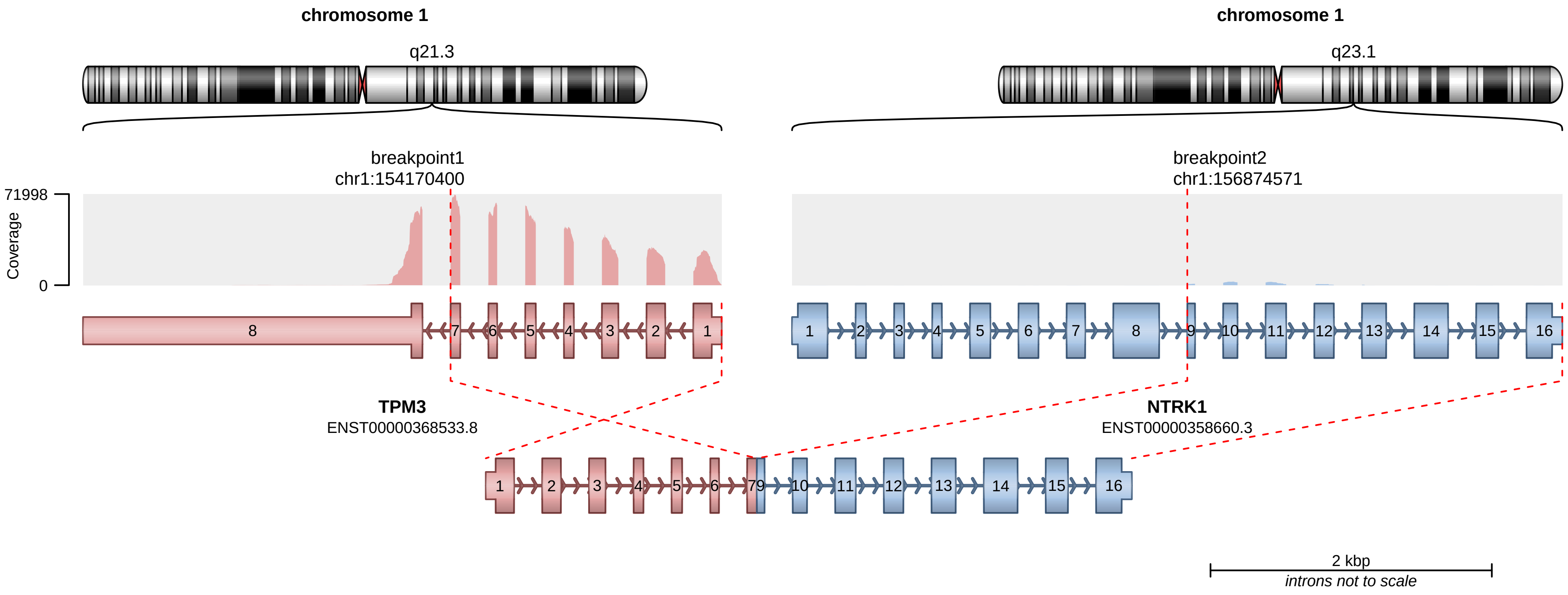


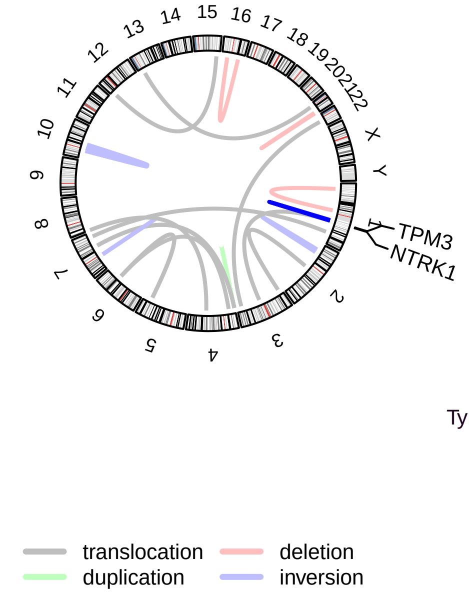
RETAINED PROTEIN DOMAINS
reading frame unclear

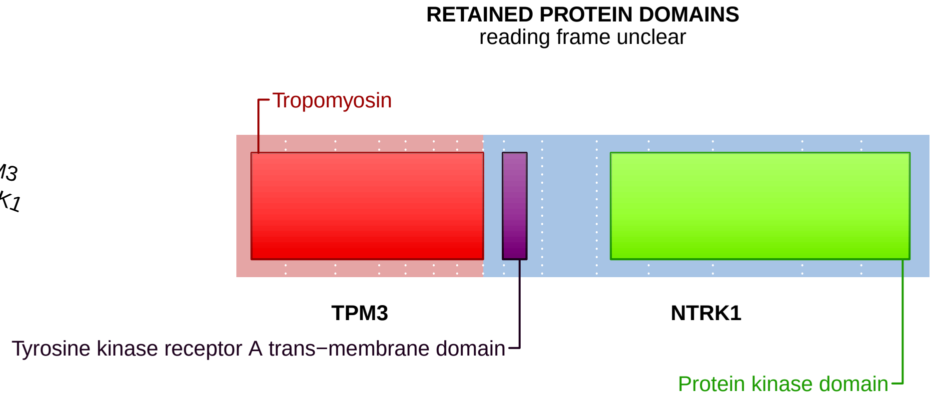


SUPPORTING READ COUNT

Split reads = 211
Discordant mates = 0

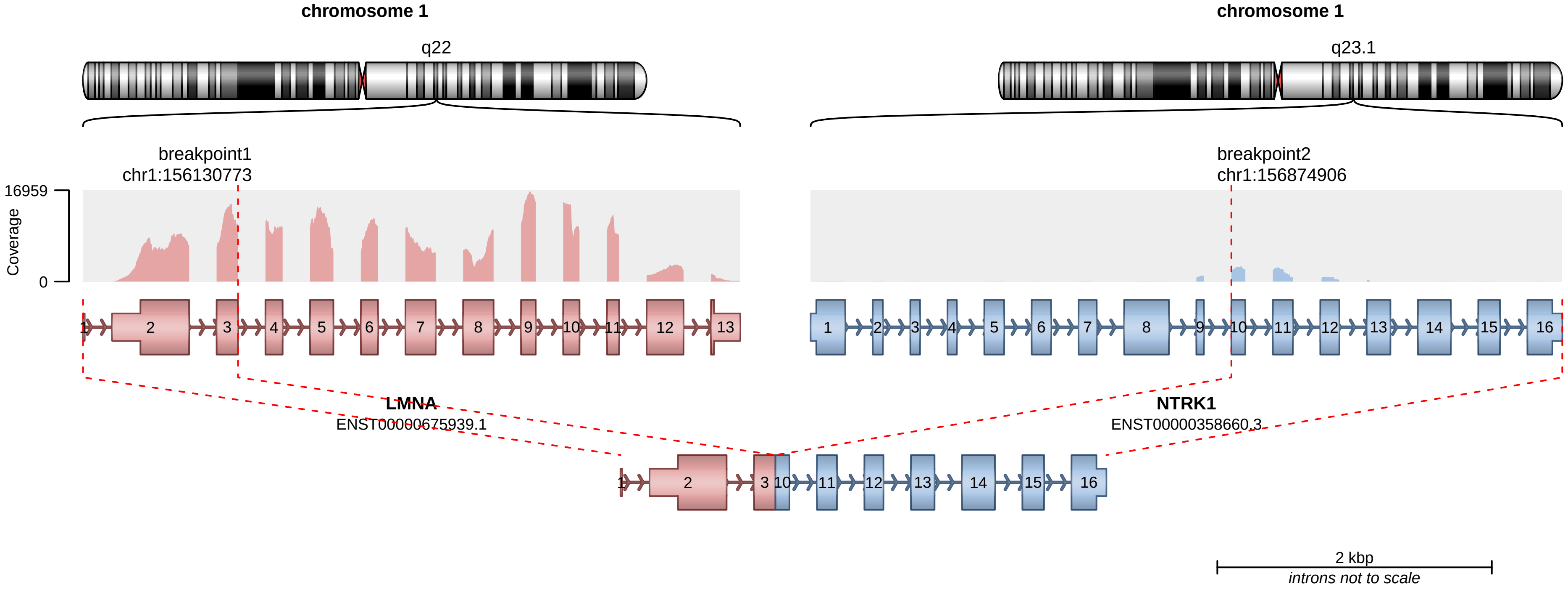


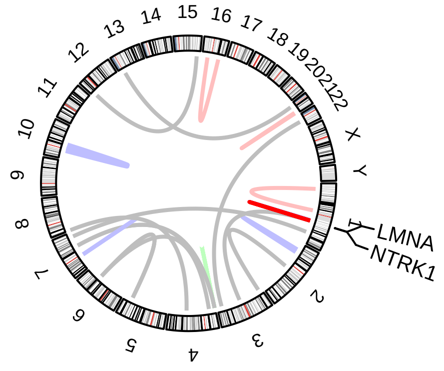




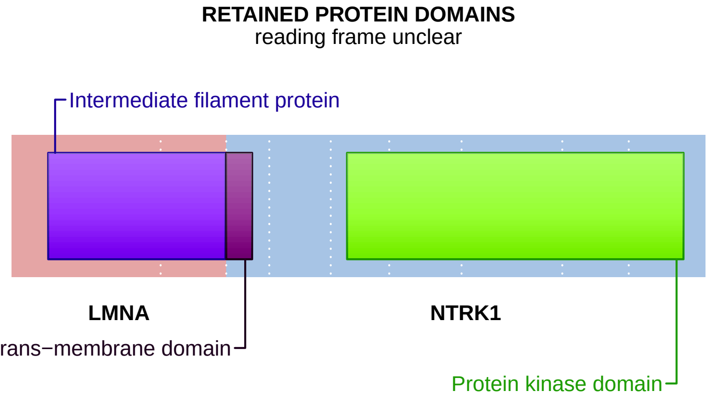
SUPPORTING READ COUNT

Split reads = 207
Discordant mates = 0



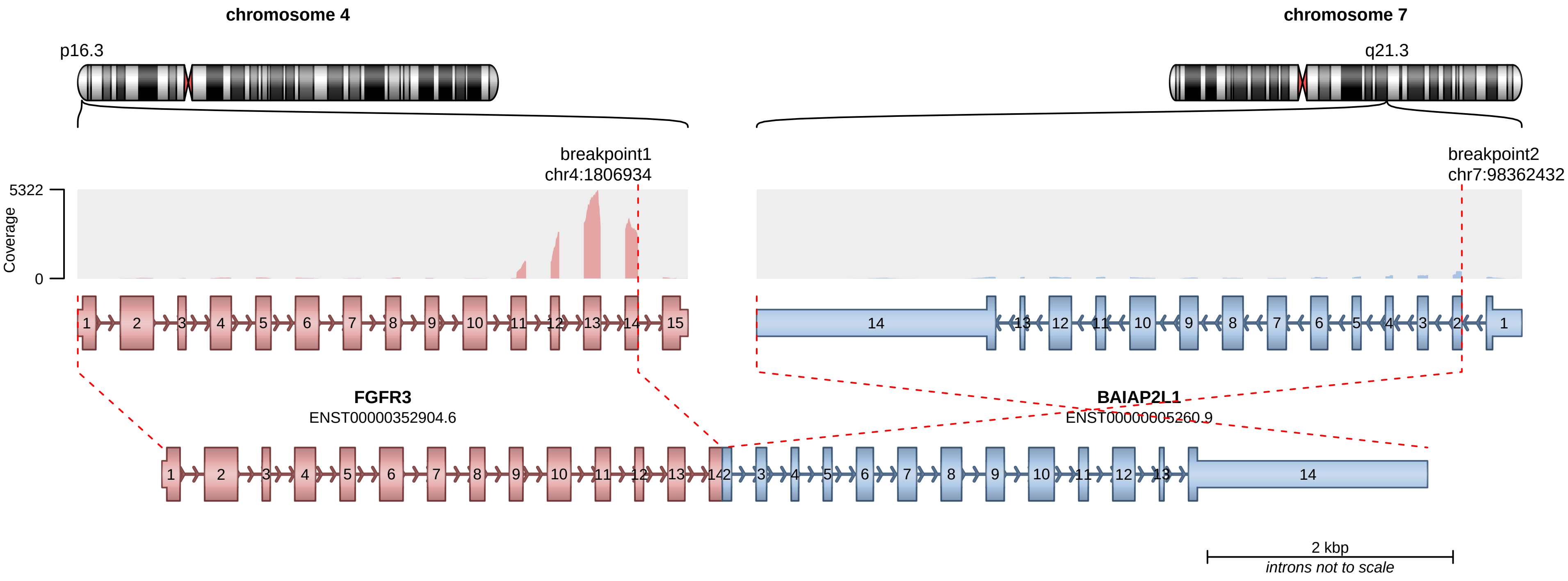


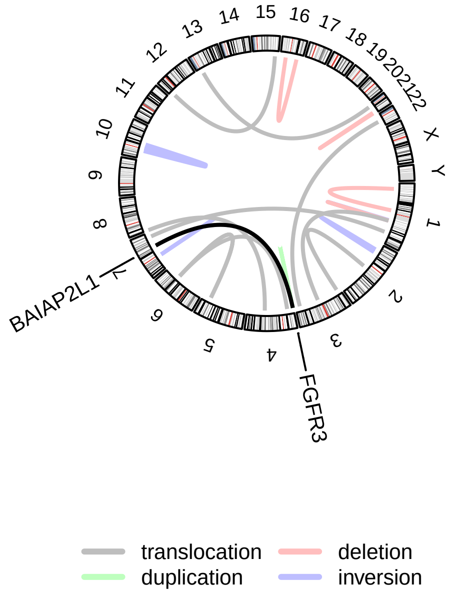
- translocation
- duplication
- deletion
- inversion



SUPPORTING READ COUNT

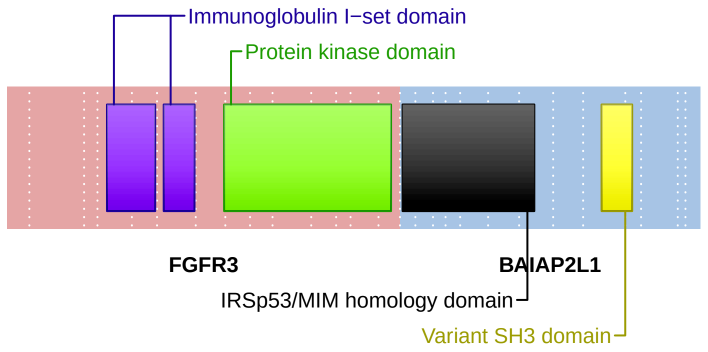
Split reads = 199
Discordant mates = 0





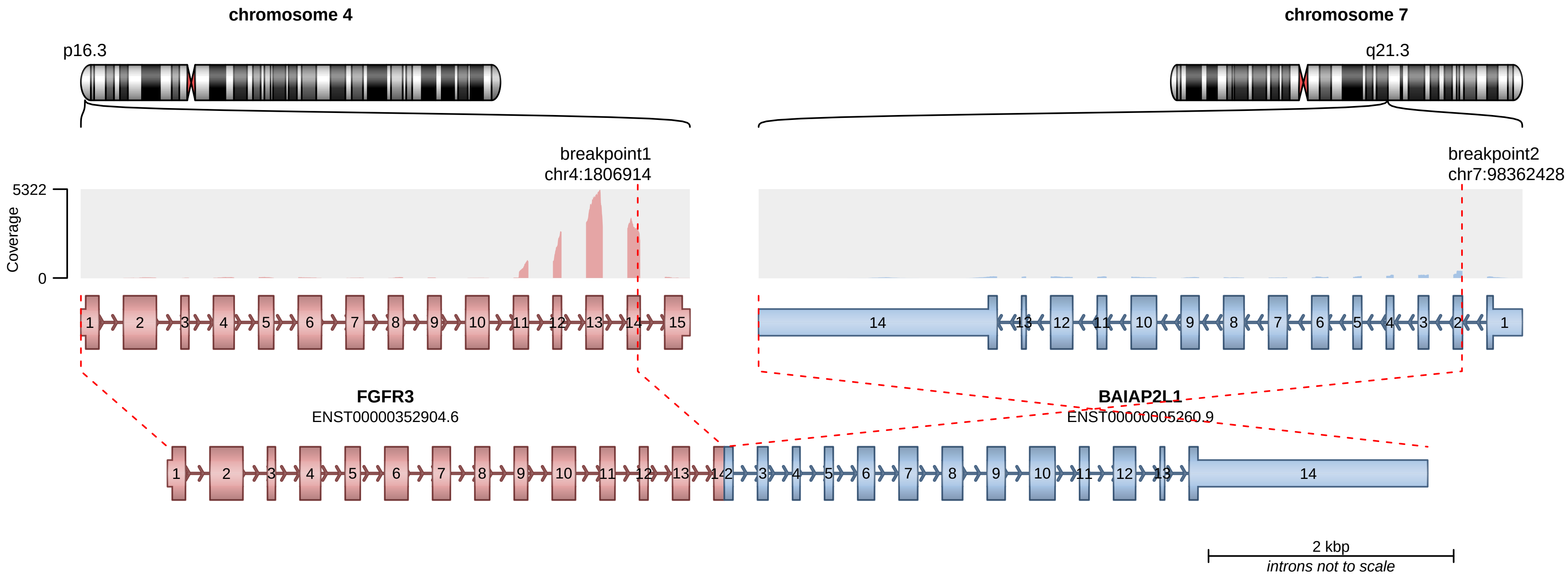
RETAINED PROTEIN DOMAINS

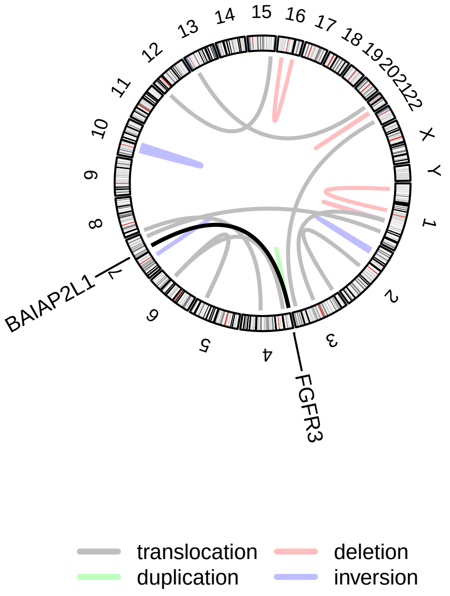
reading frame unclear

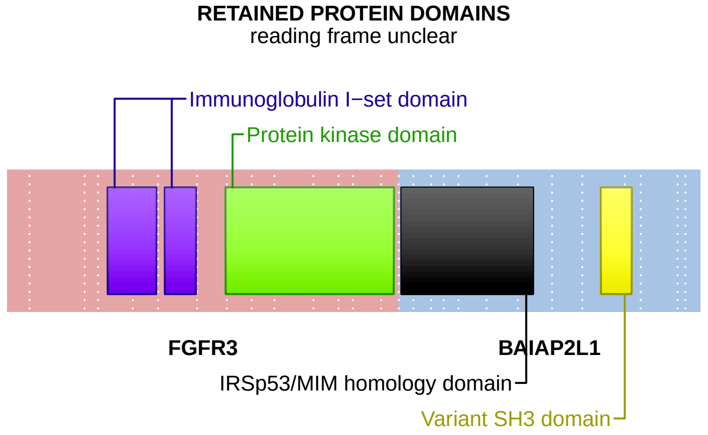


SUPPORTING READ COUNT

Split reads = 198
Discordant mates = 0

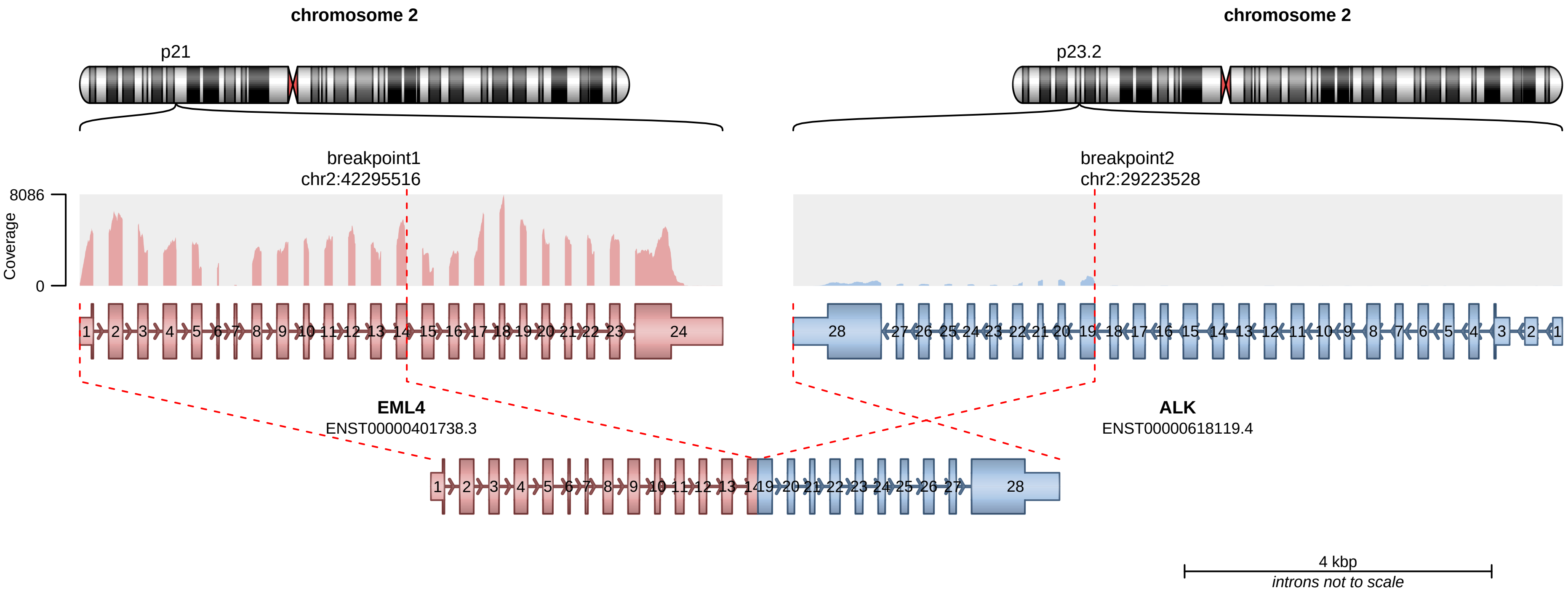


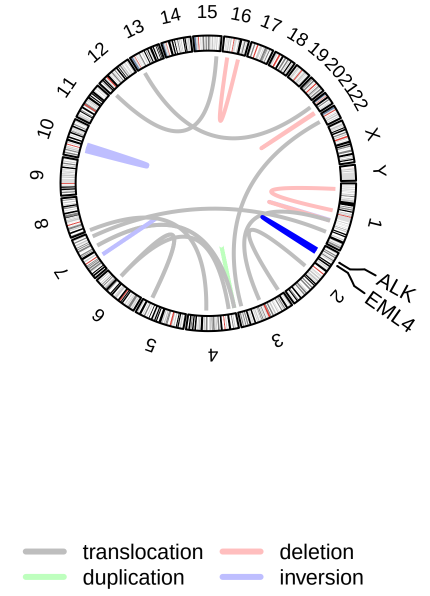


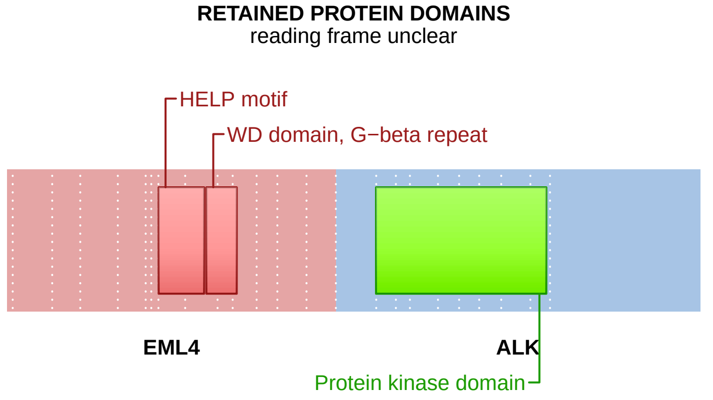


SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0

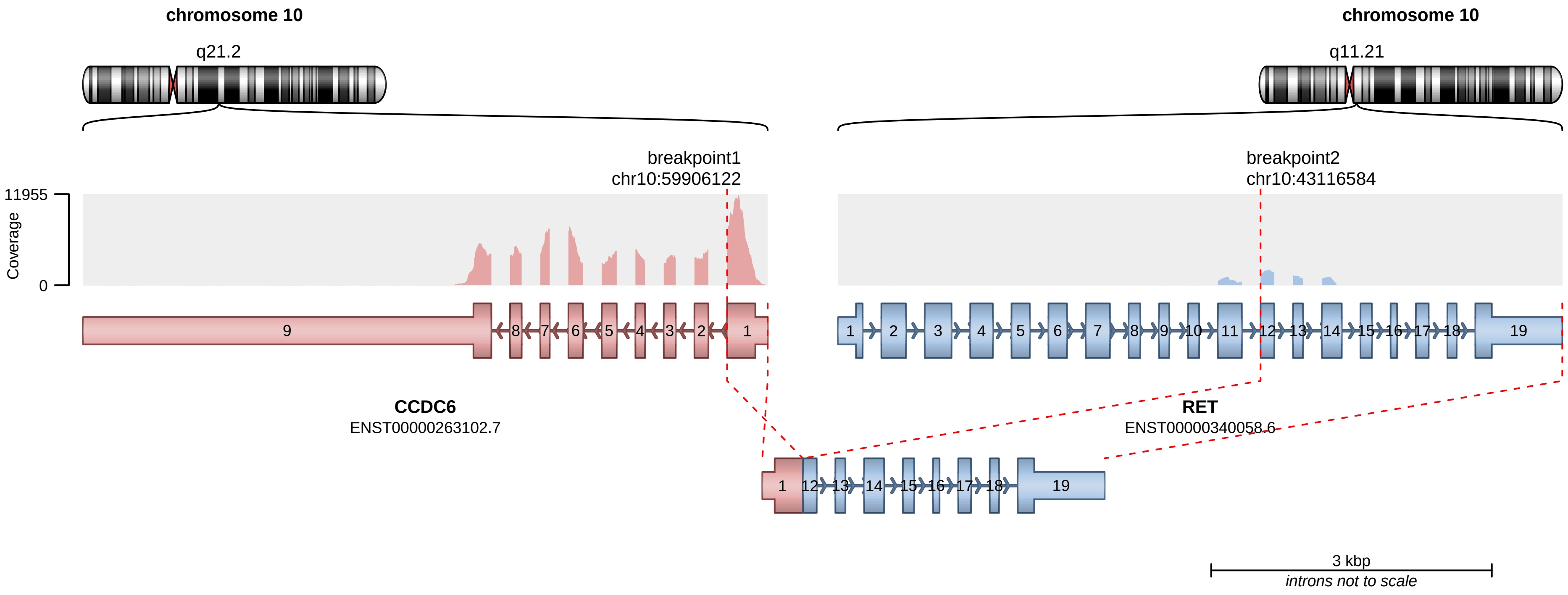


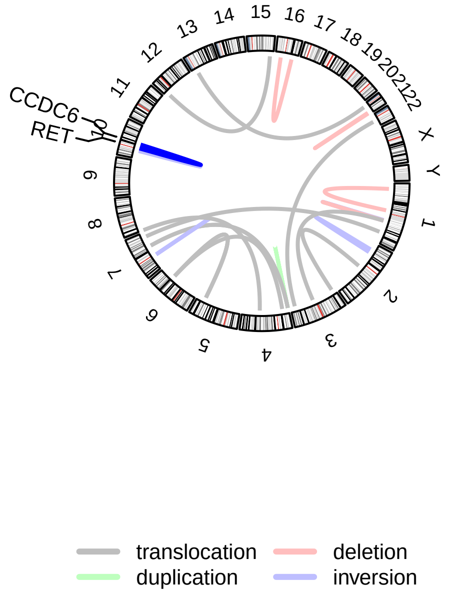


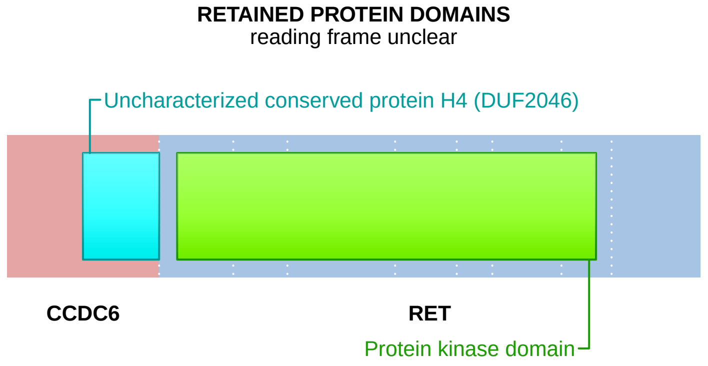


SUPPORTING READ COUNT

Split reads = 145
Discordant mates = 0

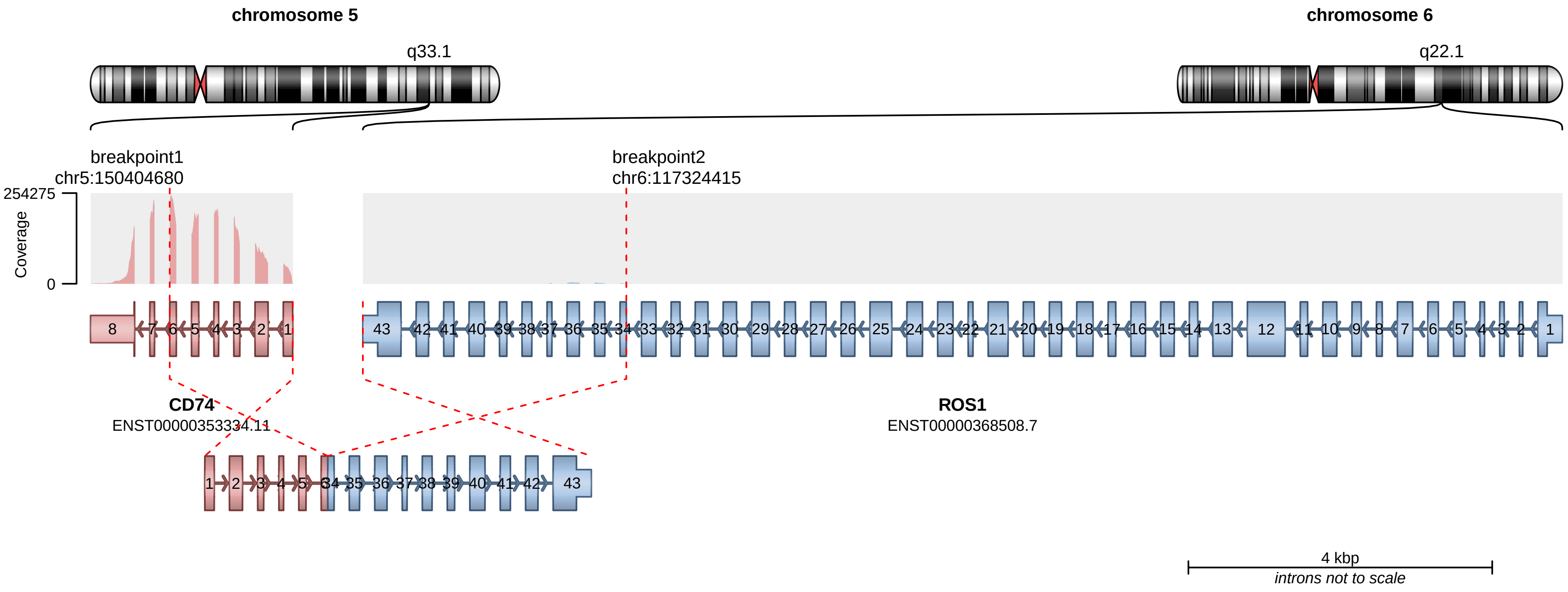


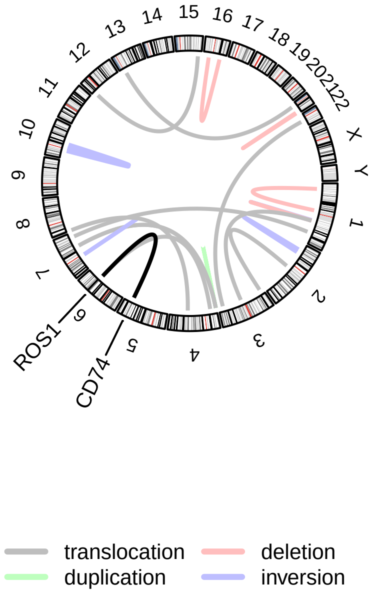


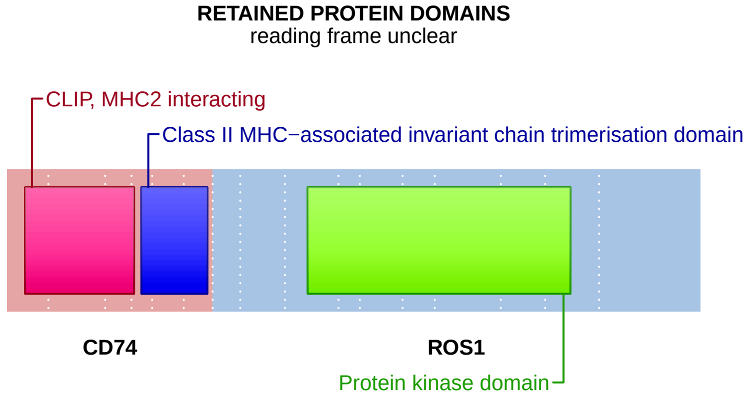


SUPPORTING READ COUNT

Split reads = 133
Discordant mates = 0

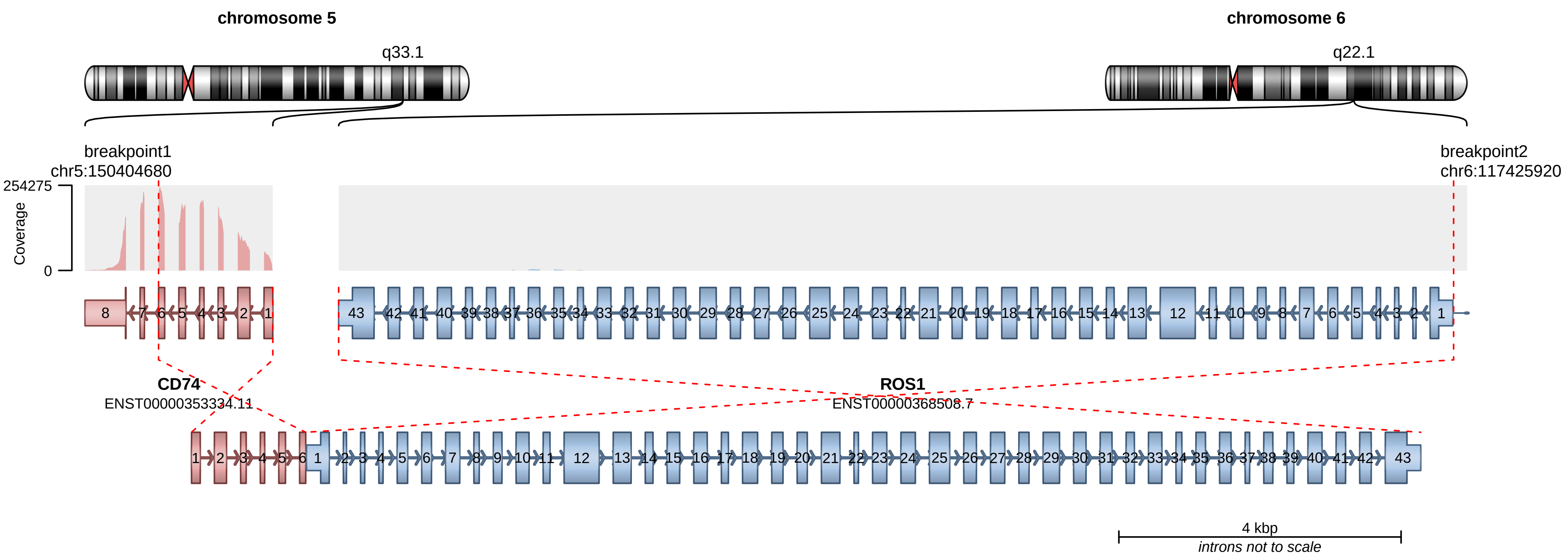


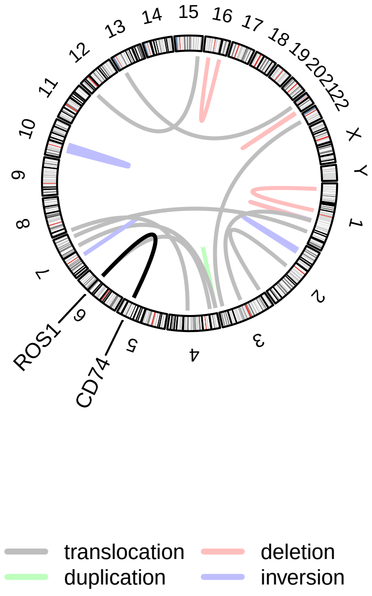


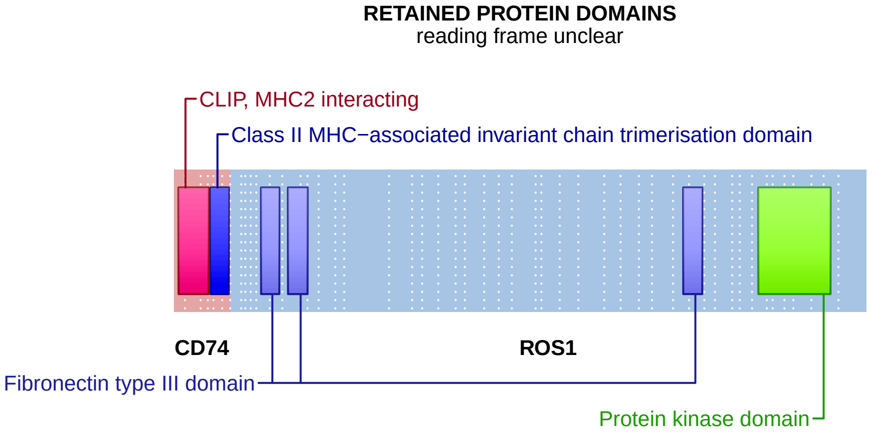


SUPPORTING READ COUNT

Split reads = 127
Discordant mates = 1

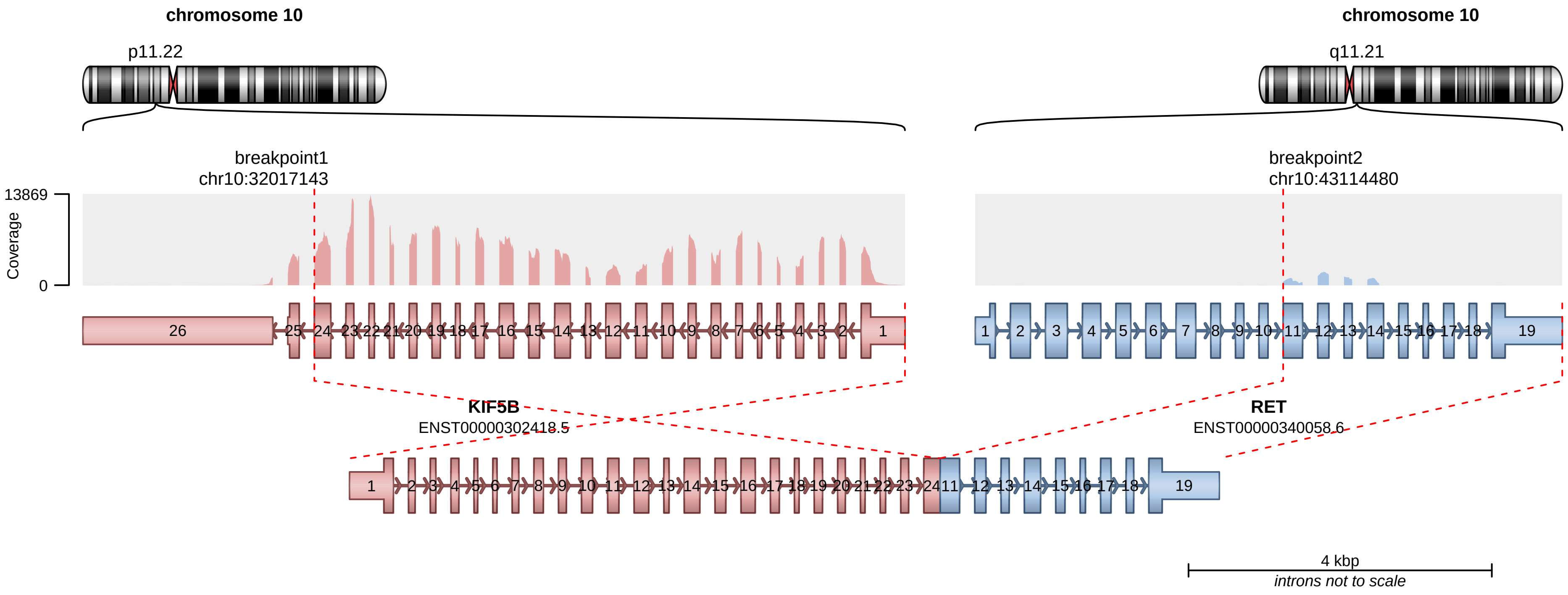


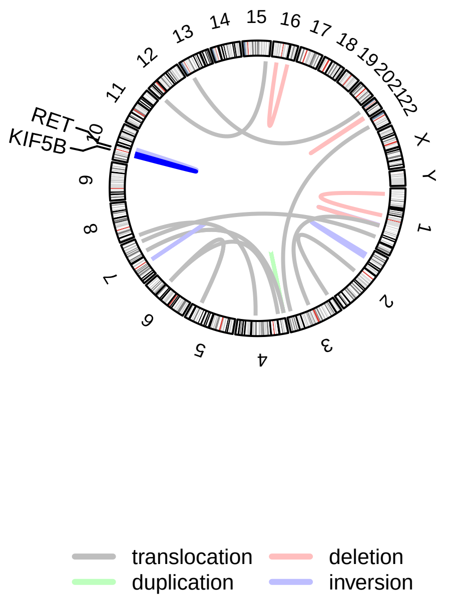




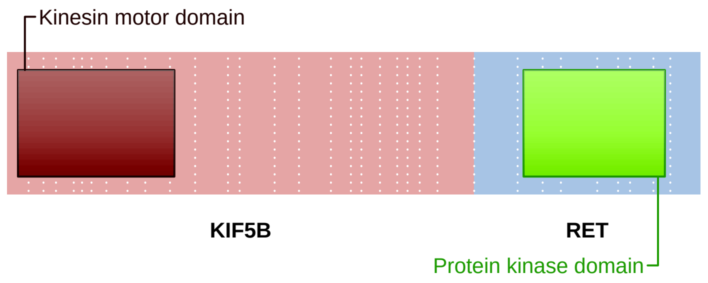
SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 1



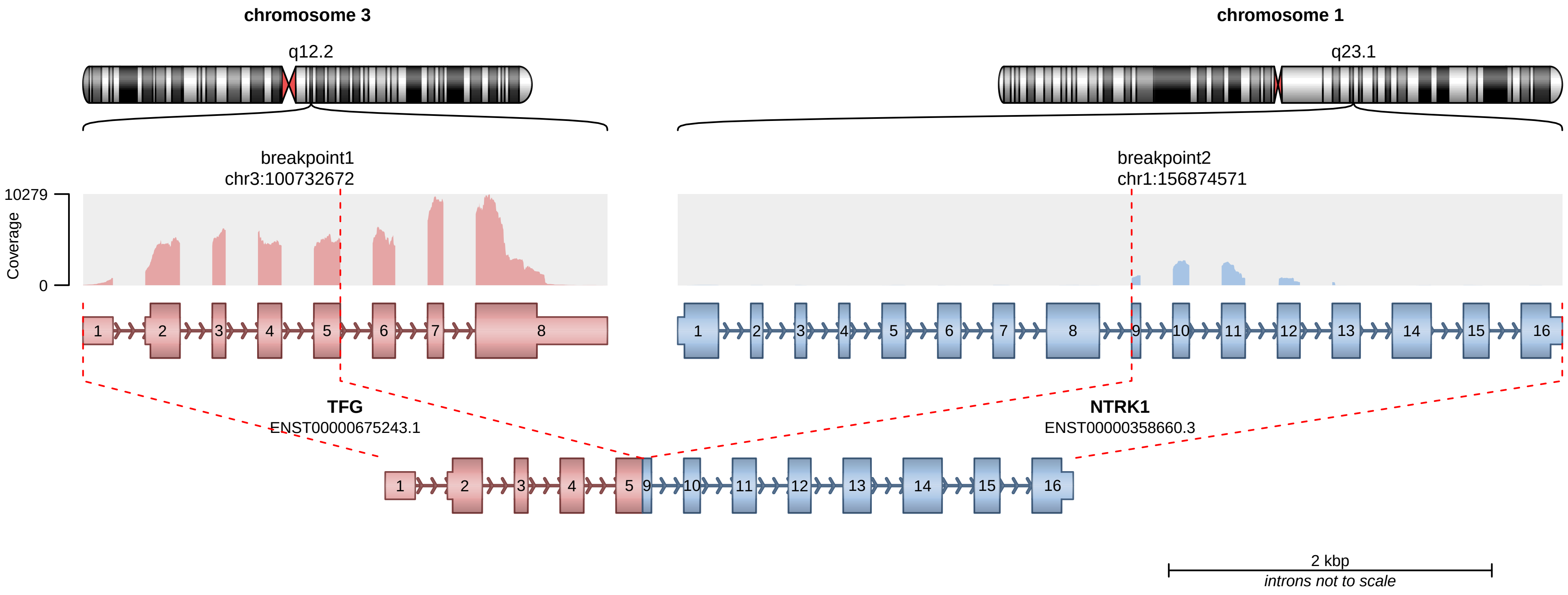


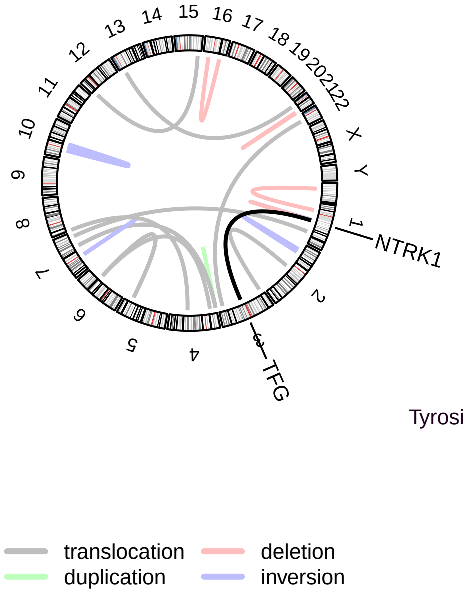
RETAINED PROTEIN DOMAINS
reading frame unclear



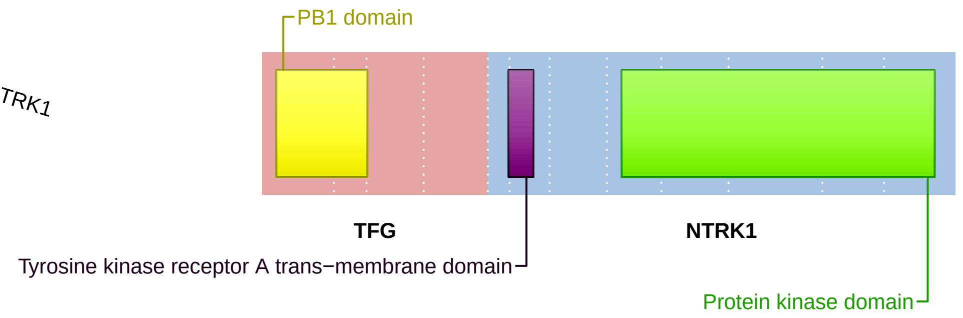
SUPPORTING READ COUNT

Split reads = 127
Discordant mates = 1



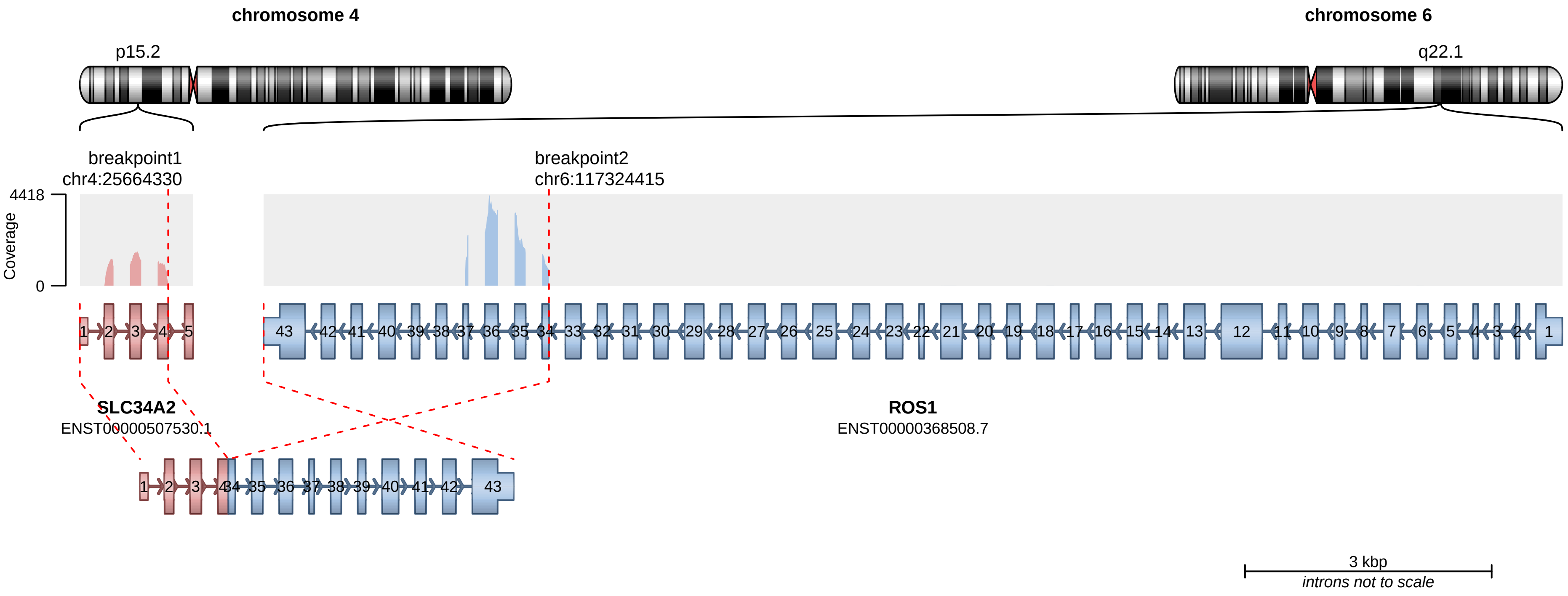


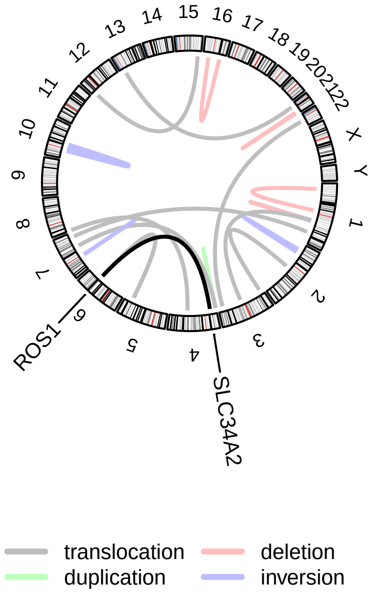
RETAINED PROTEIN DOMAINS
reading frame unclear

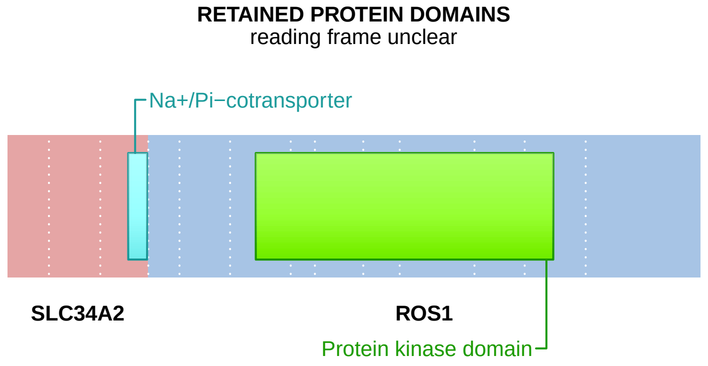


SUPPORTING READ COUNT

Split reads = 79
Discordant mates = 1

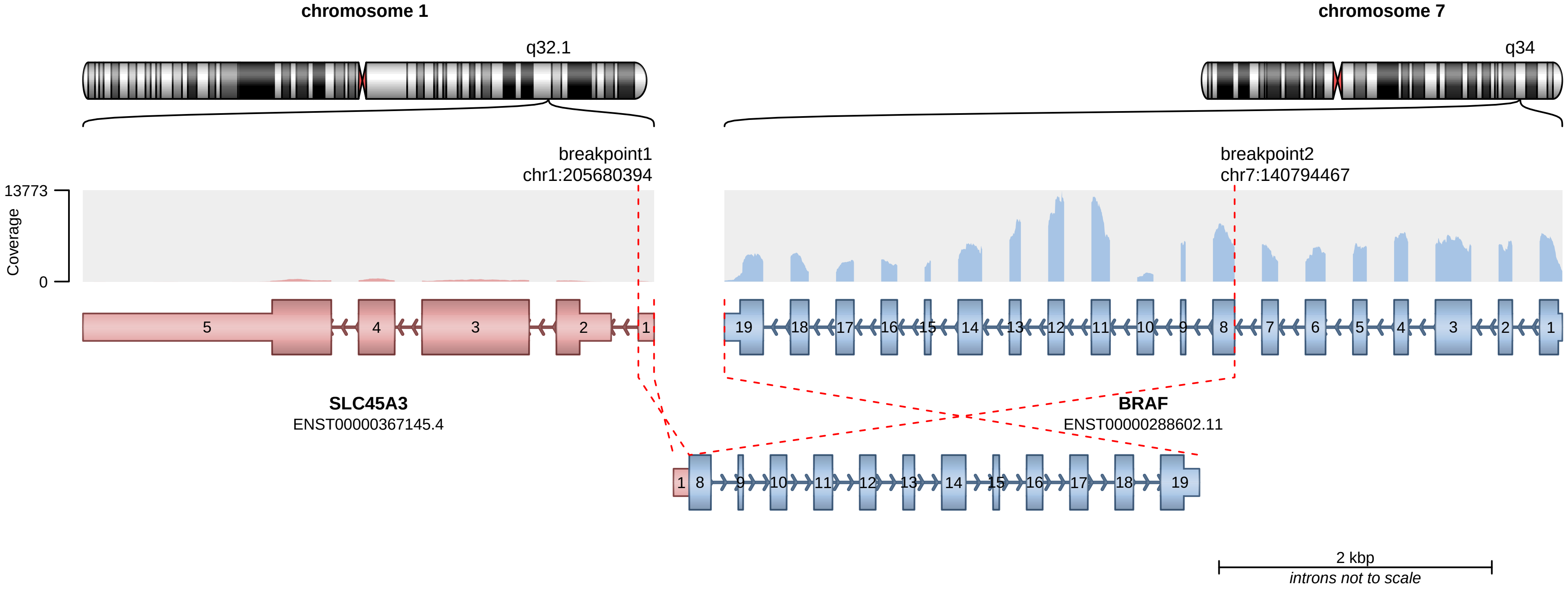


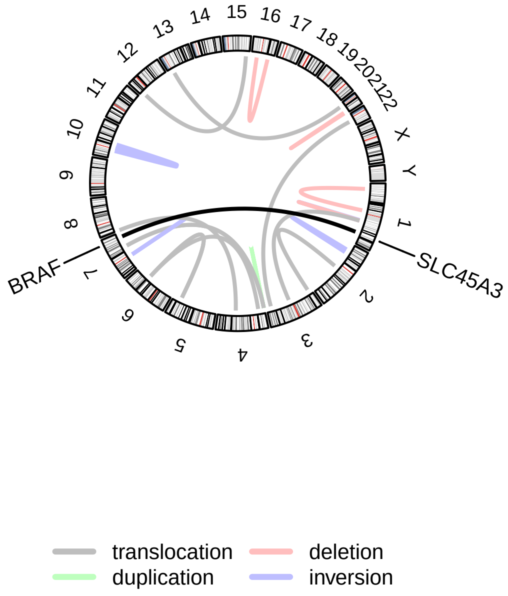




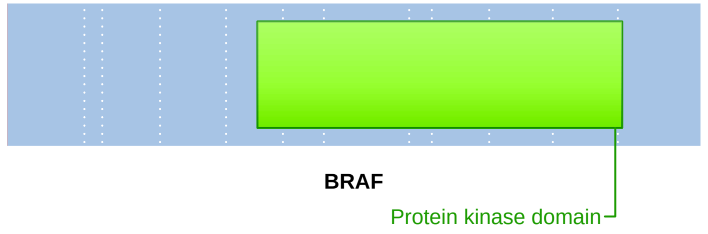
SUPPORTING READ COUNT

Split reads = 75
Discordant mates = 0



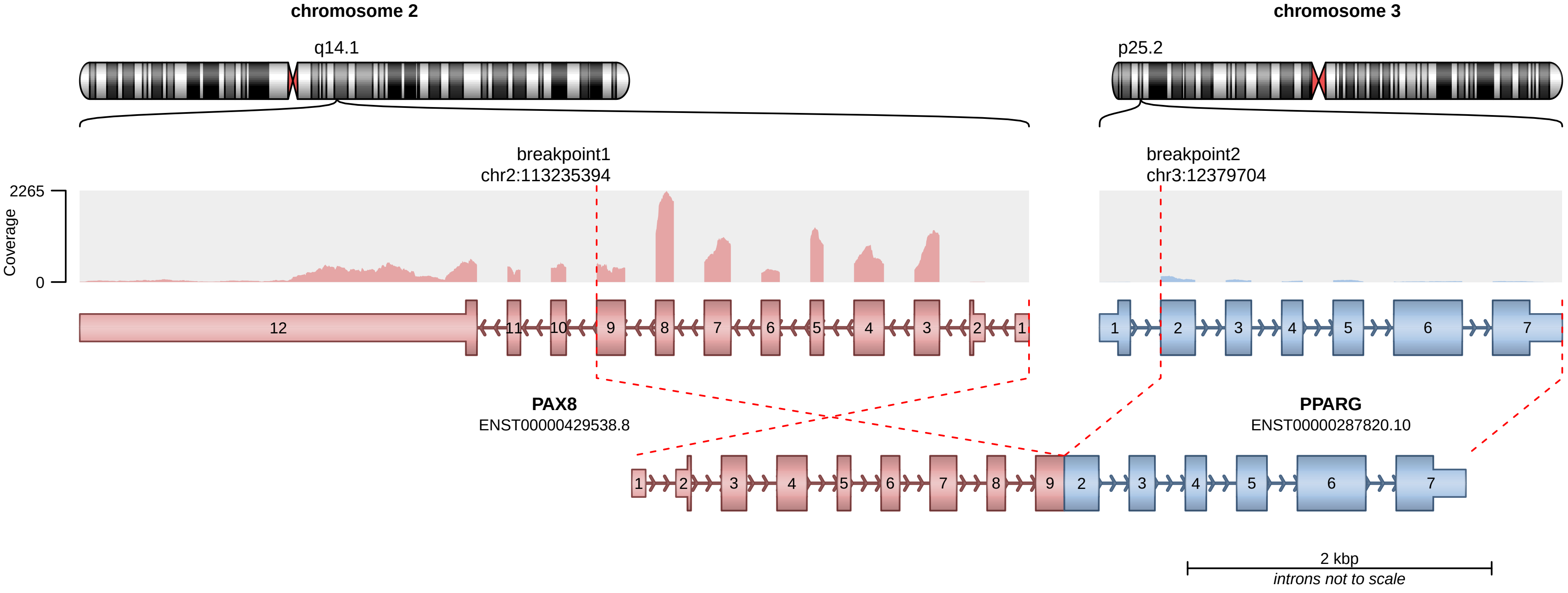


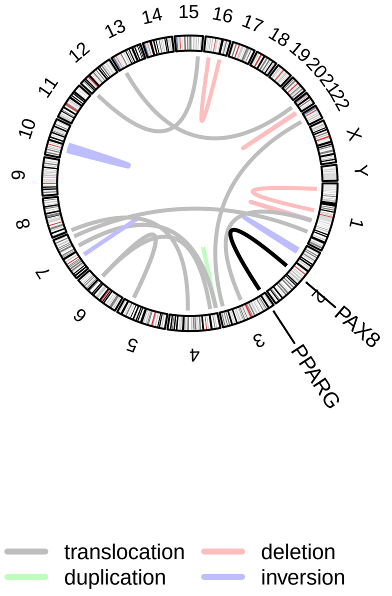
RETAINED PROTEIN DOMAINS
reading frame unclear



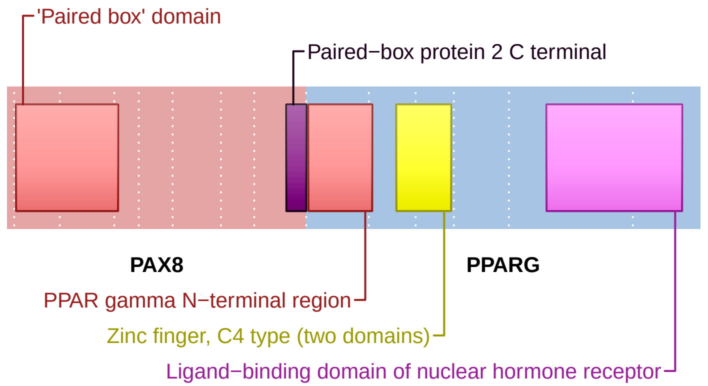
SUPPORTING READ COUNT

Split reads = 65
Discordant mates = 0



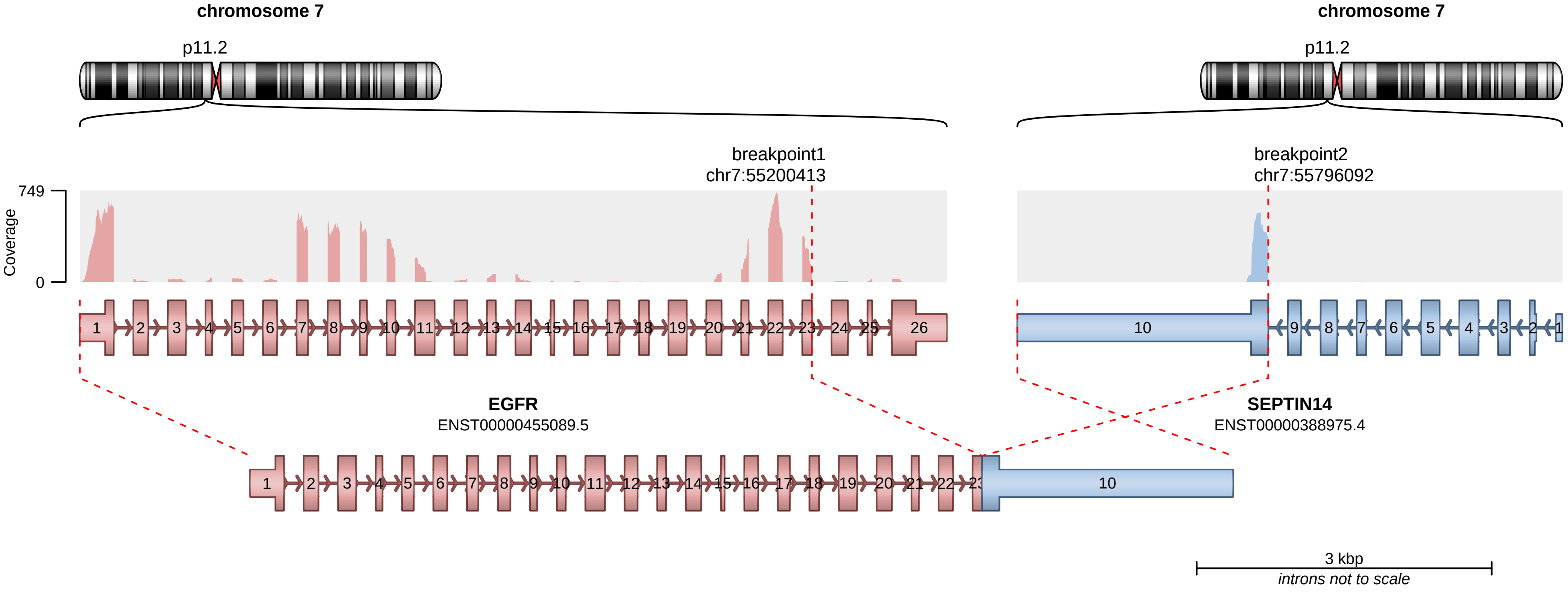


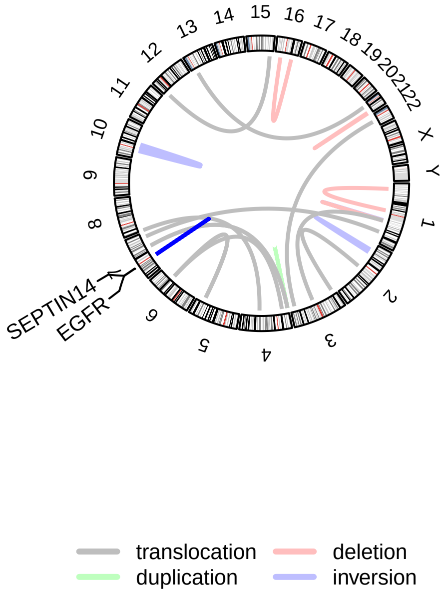
RETAINED PROTEIN DOMAINS
reading frame unclear



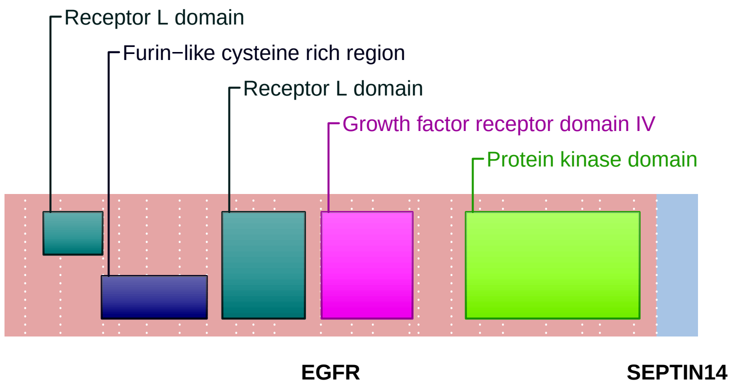
SUPPORTING READ COUNT

Split reads = 46
Discordant mates = 0



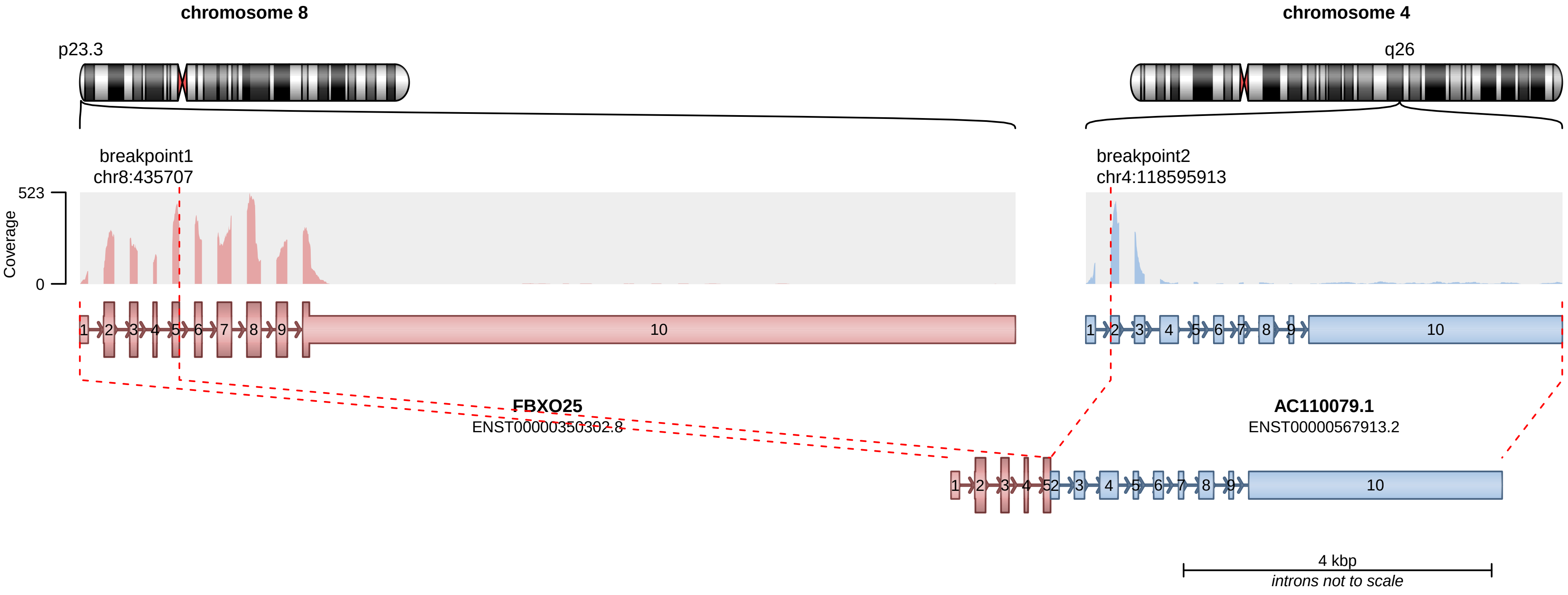


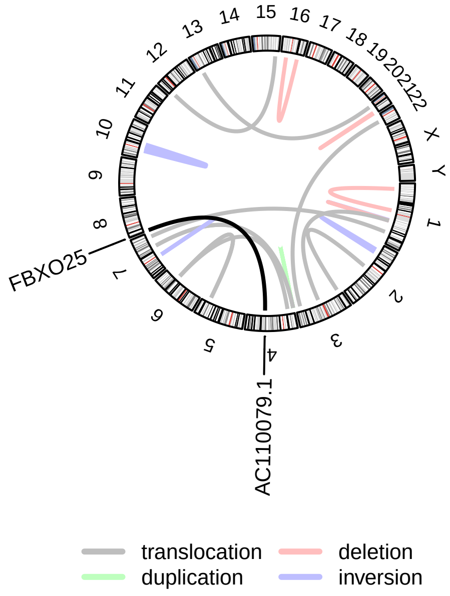
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 43
Discordant mates = 0

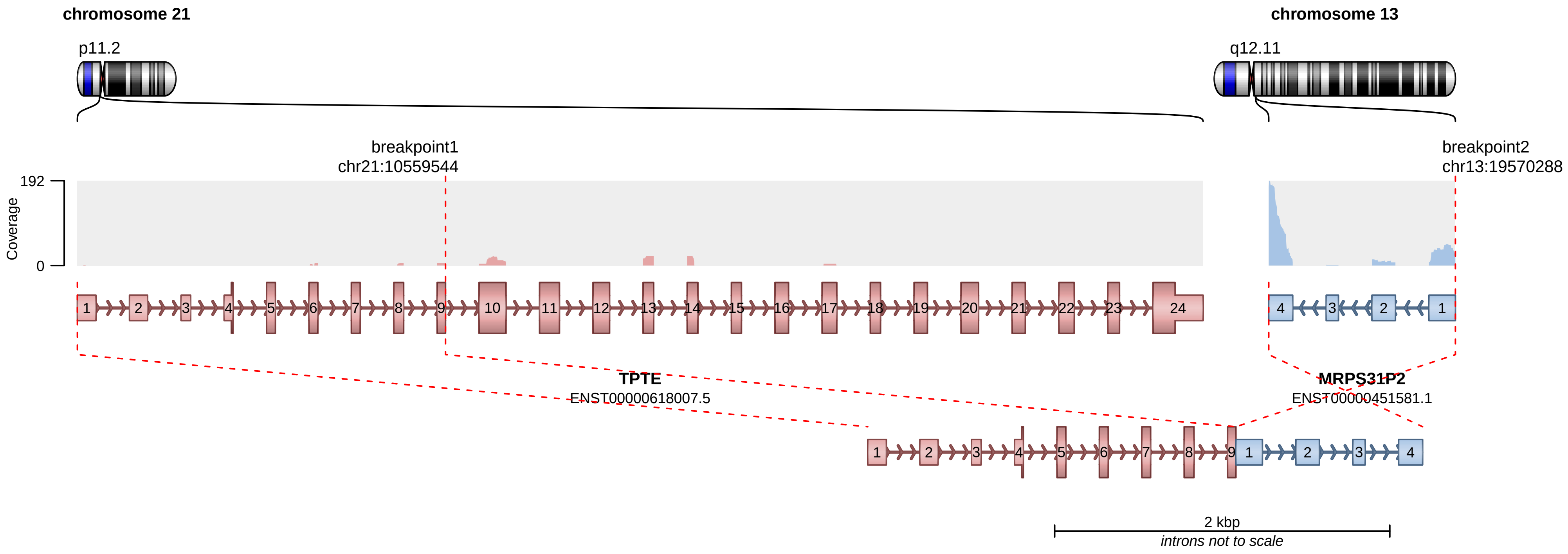


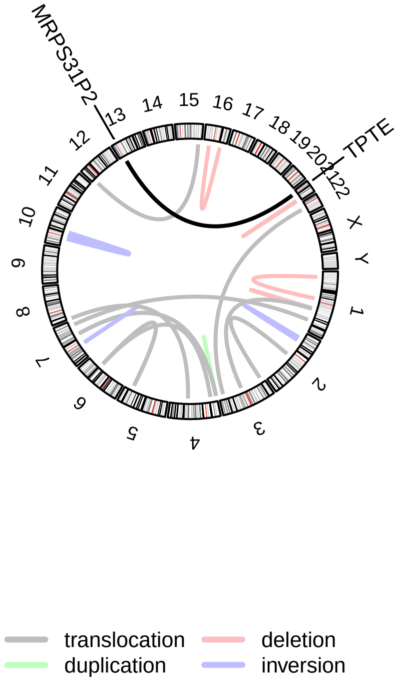


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 5
Discordant mates = 0

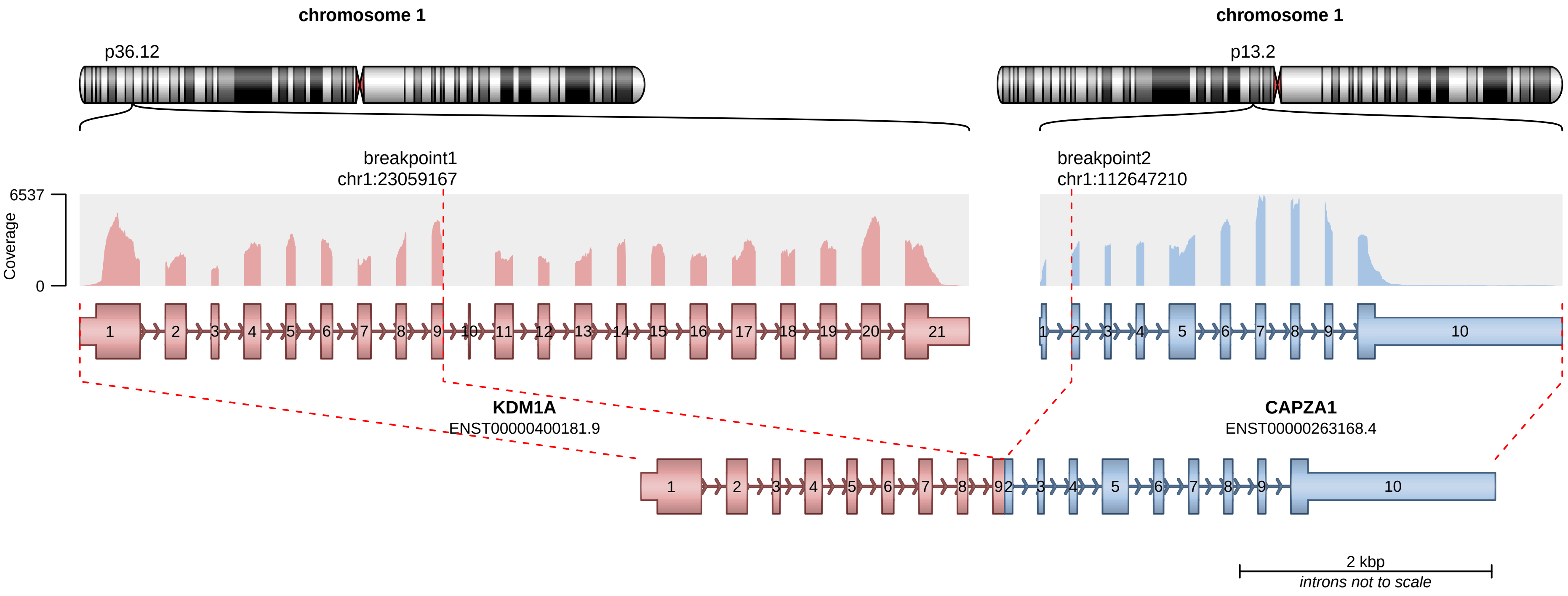


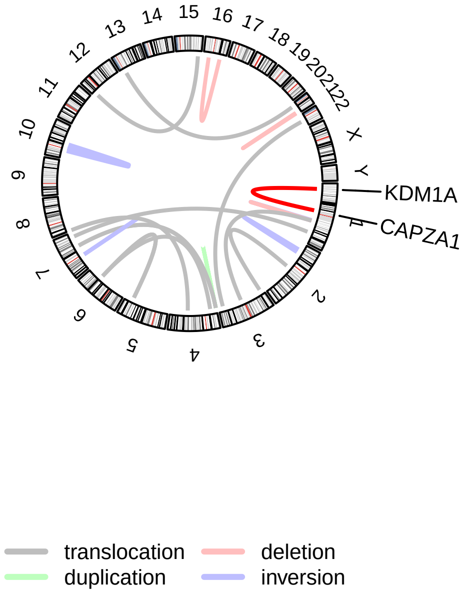


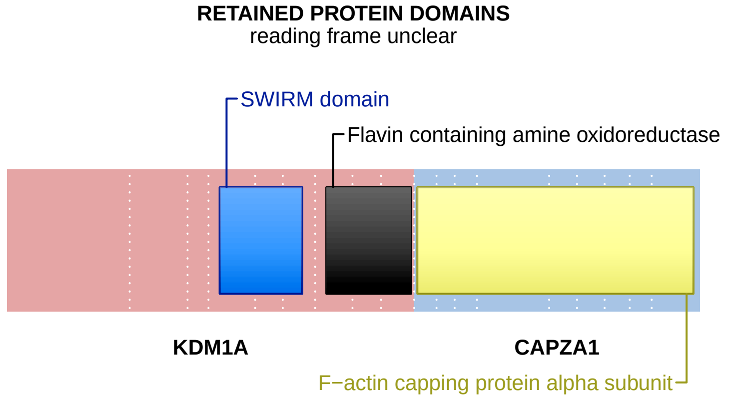
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0

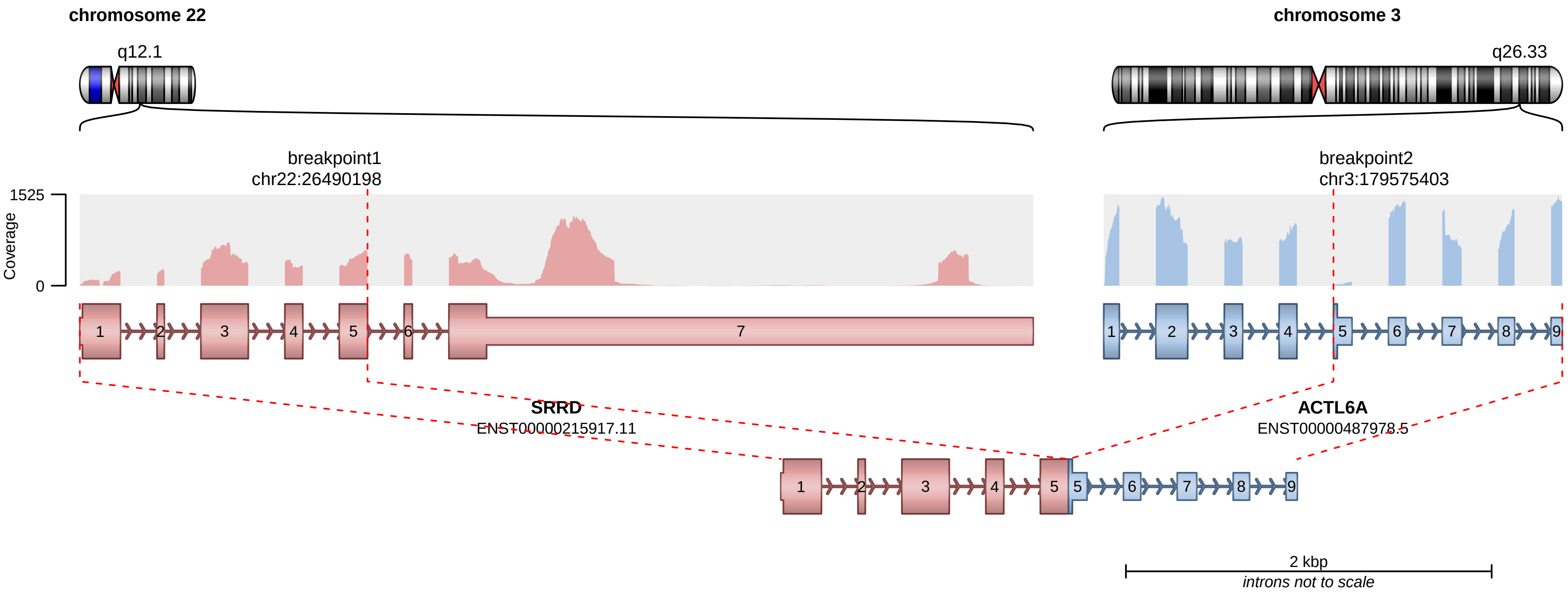


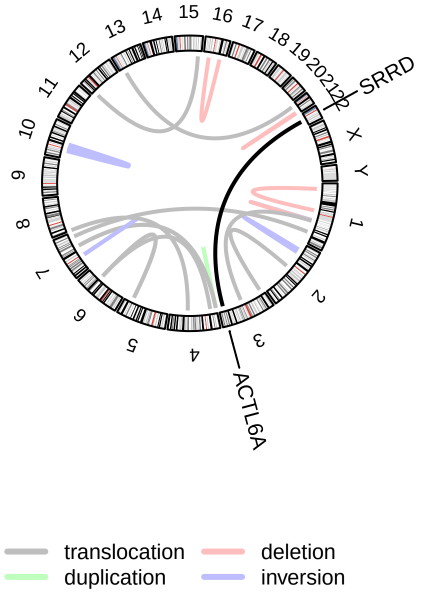


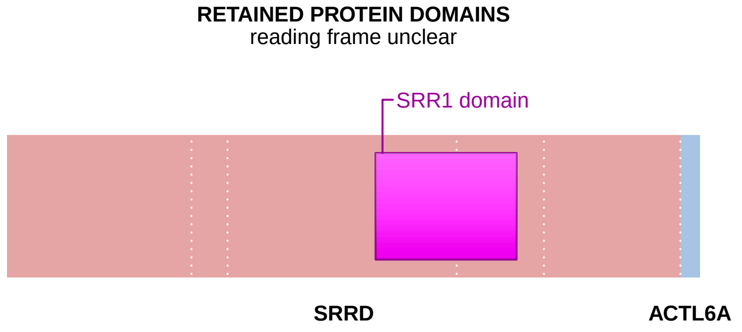


SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

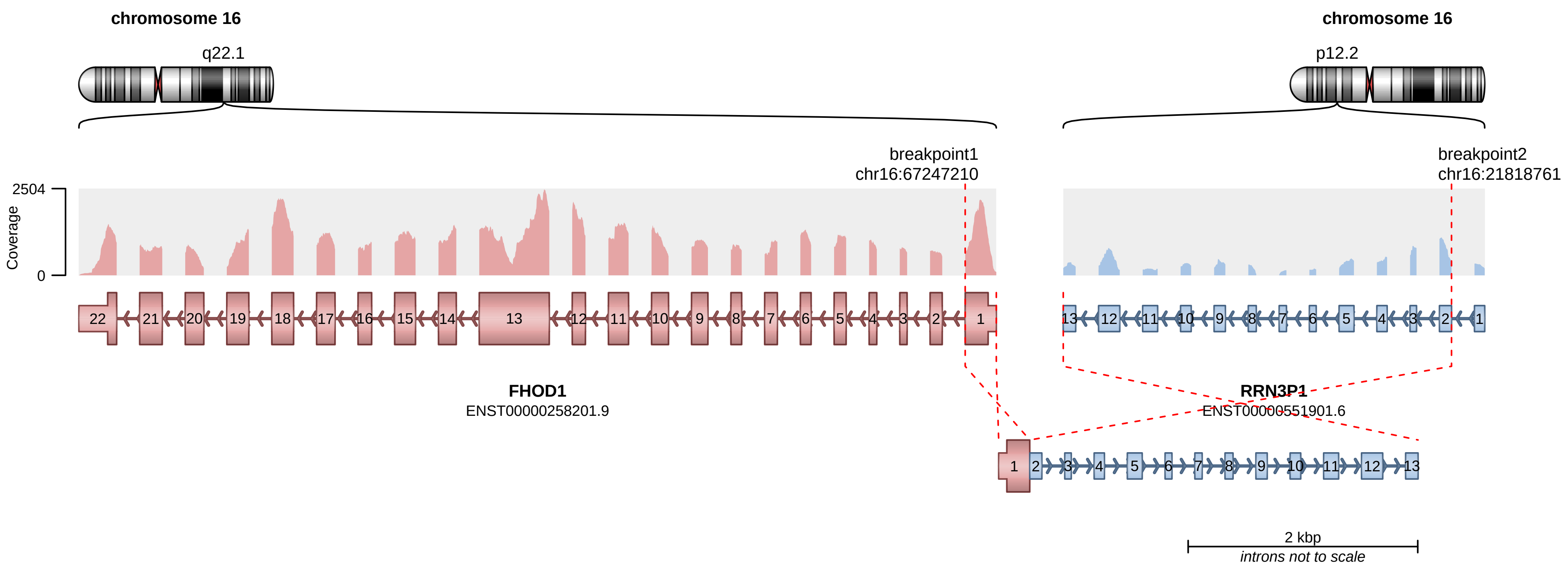


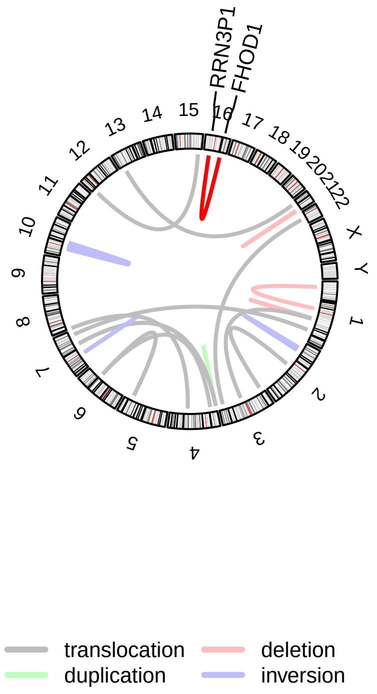


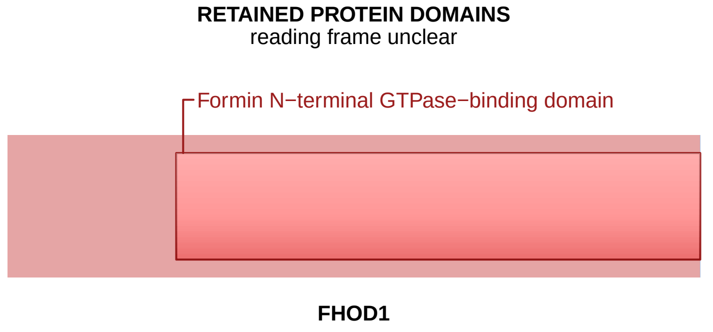


SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0







SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0