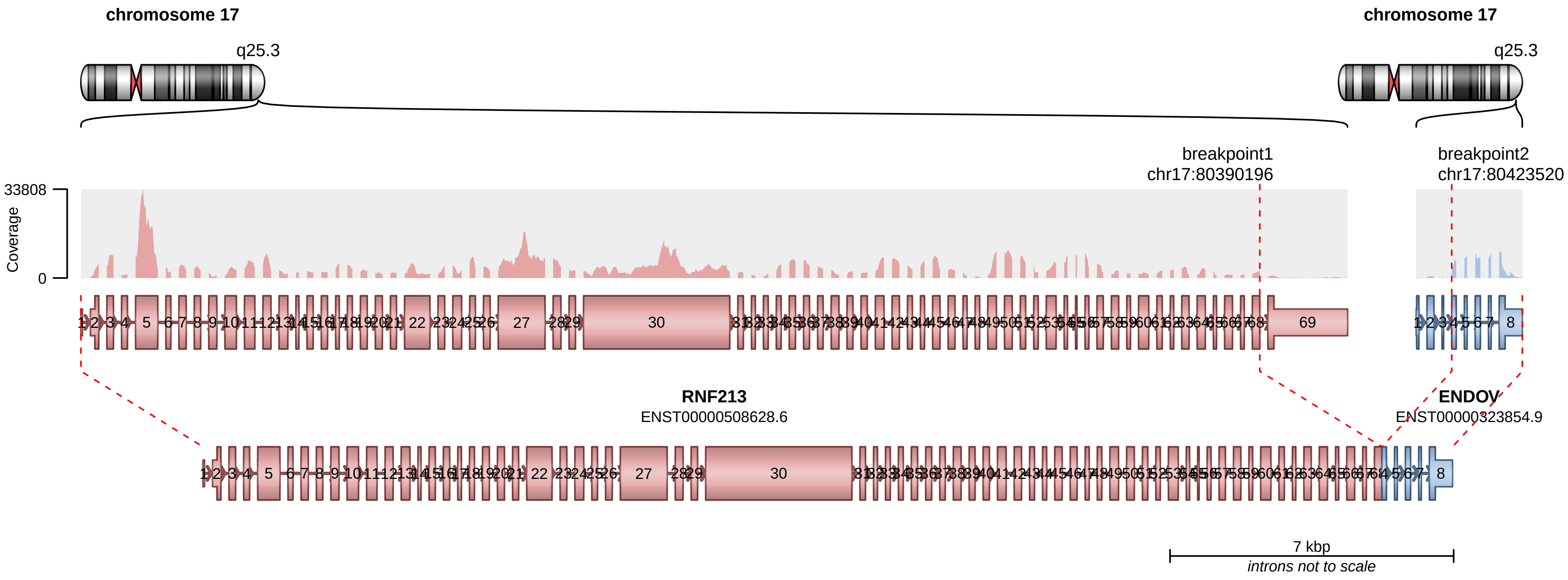
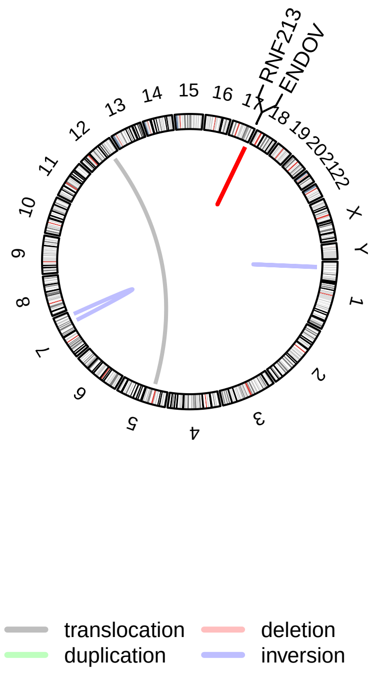
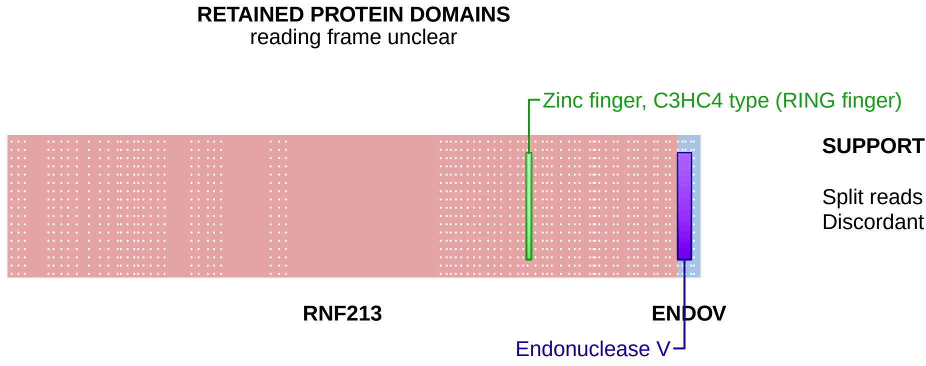
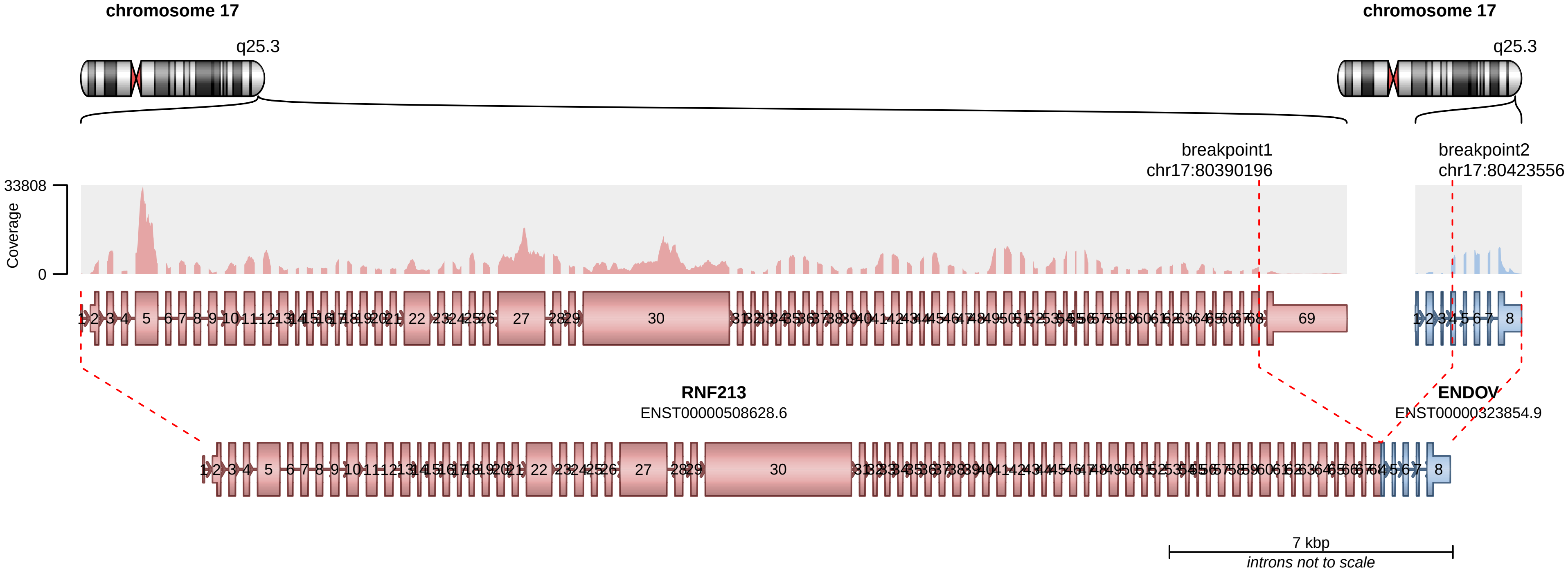


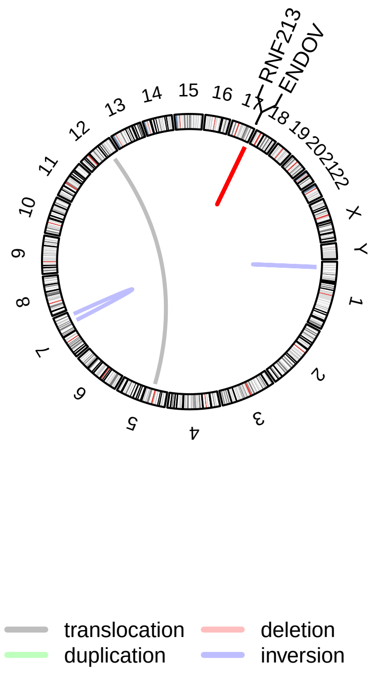




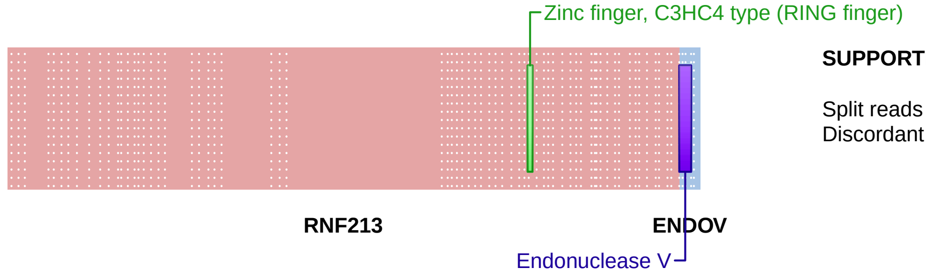
SUPPORTING READ COUNT

Split reads = 681
Discordant mates = 6



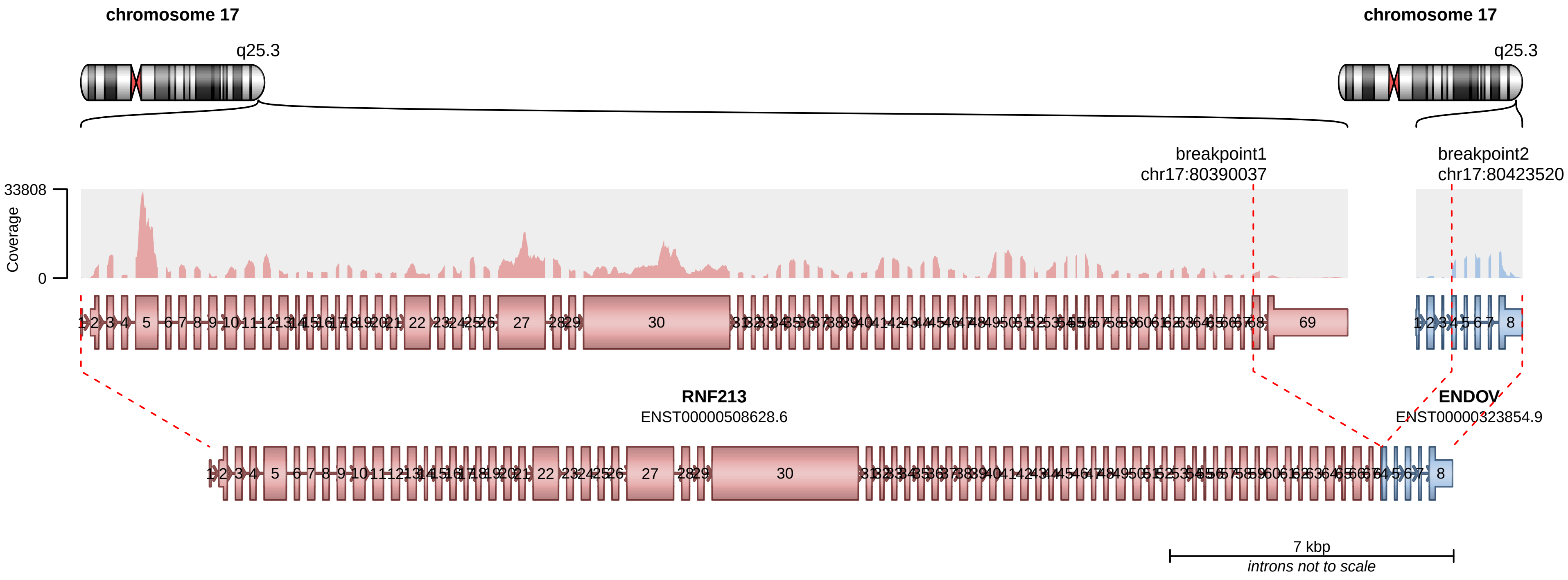


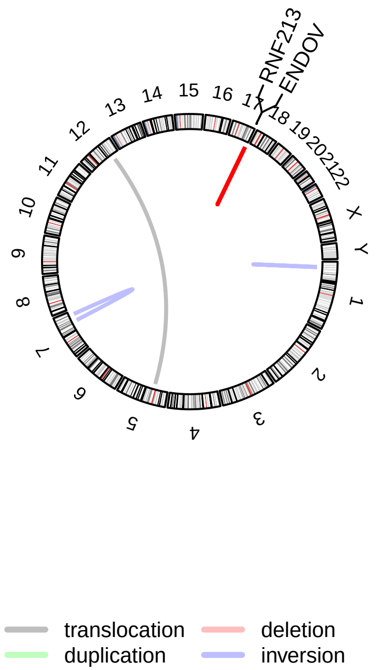
RETAINED PROTEIN DOMAINS
reading frame unclear

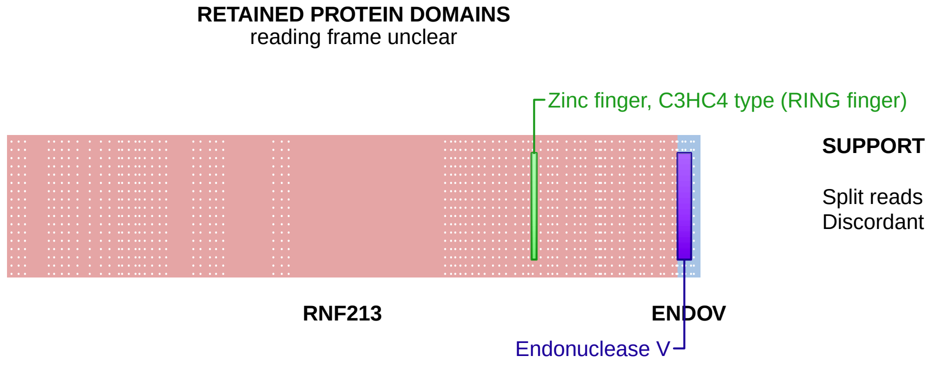


SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 4

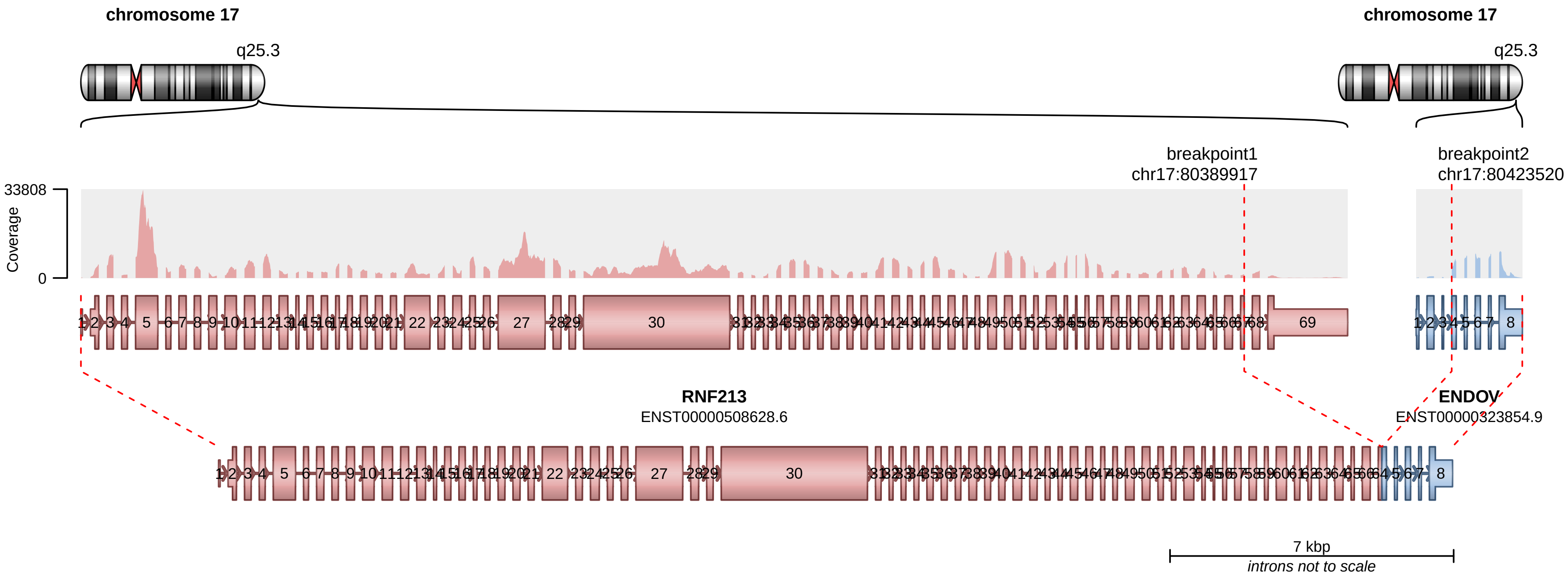


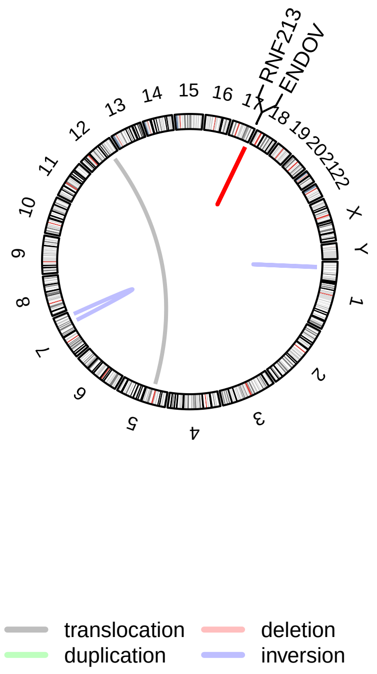




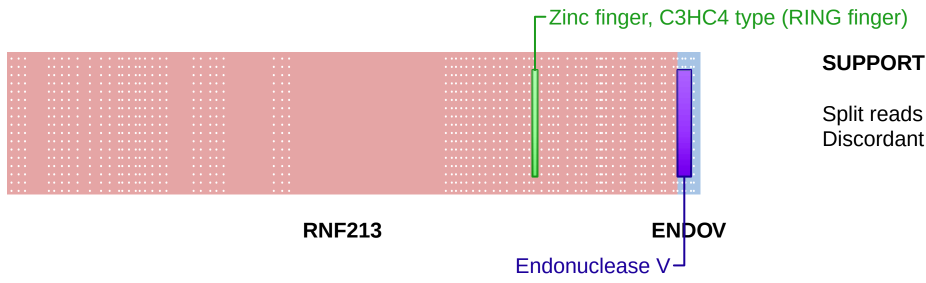
SUPPORTING READ COUNT

Split reads = 4
Discordant mates = 0



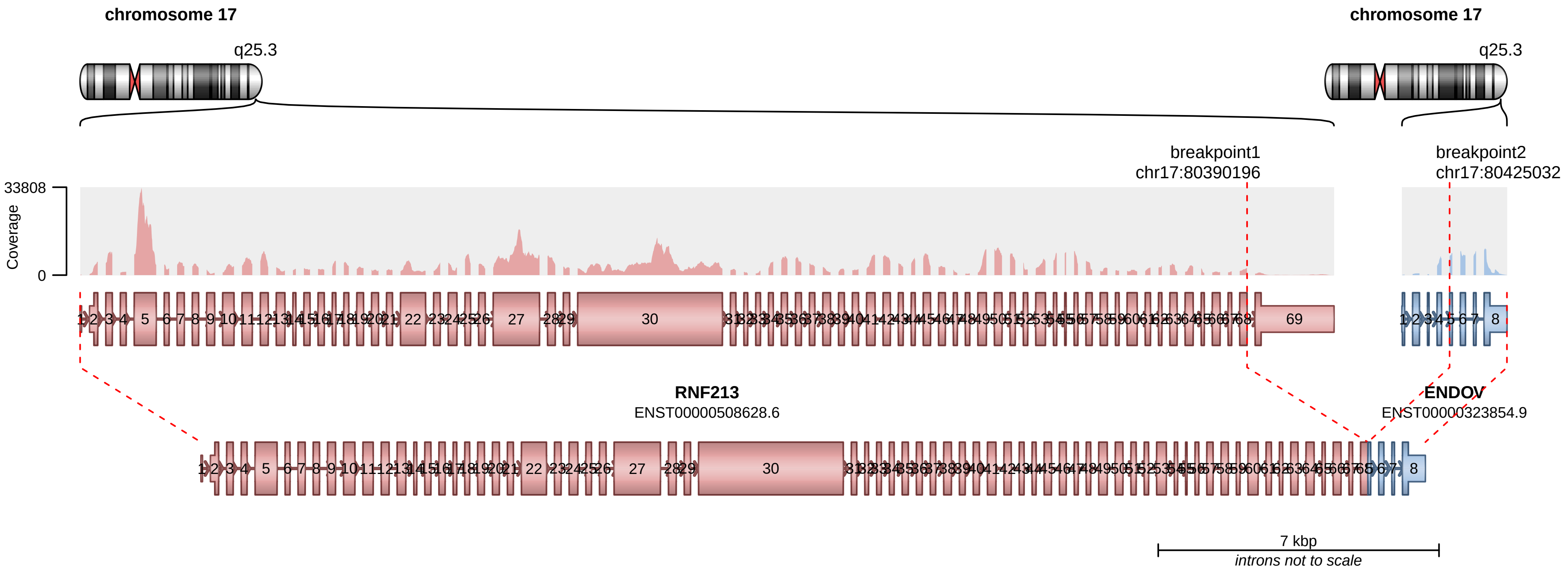


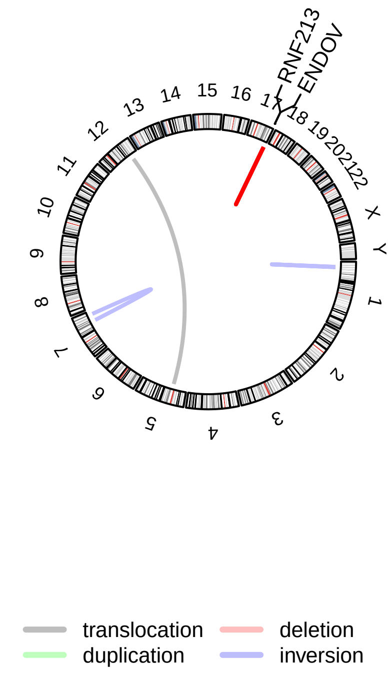
RETAINED PROTEIN DOMAINS
reading frame unclear



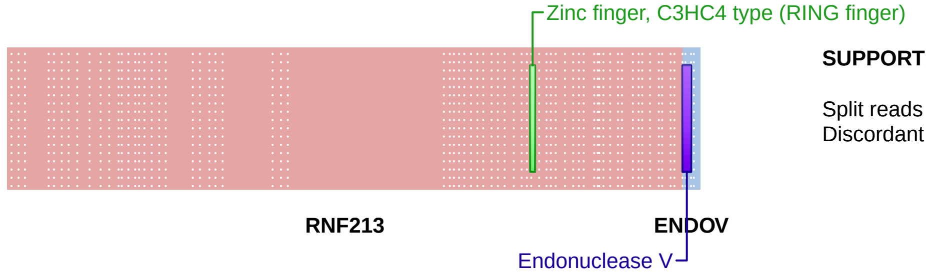
SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0



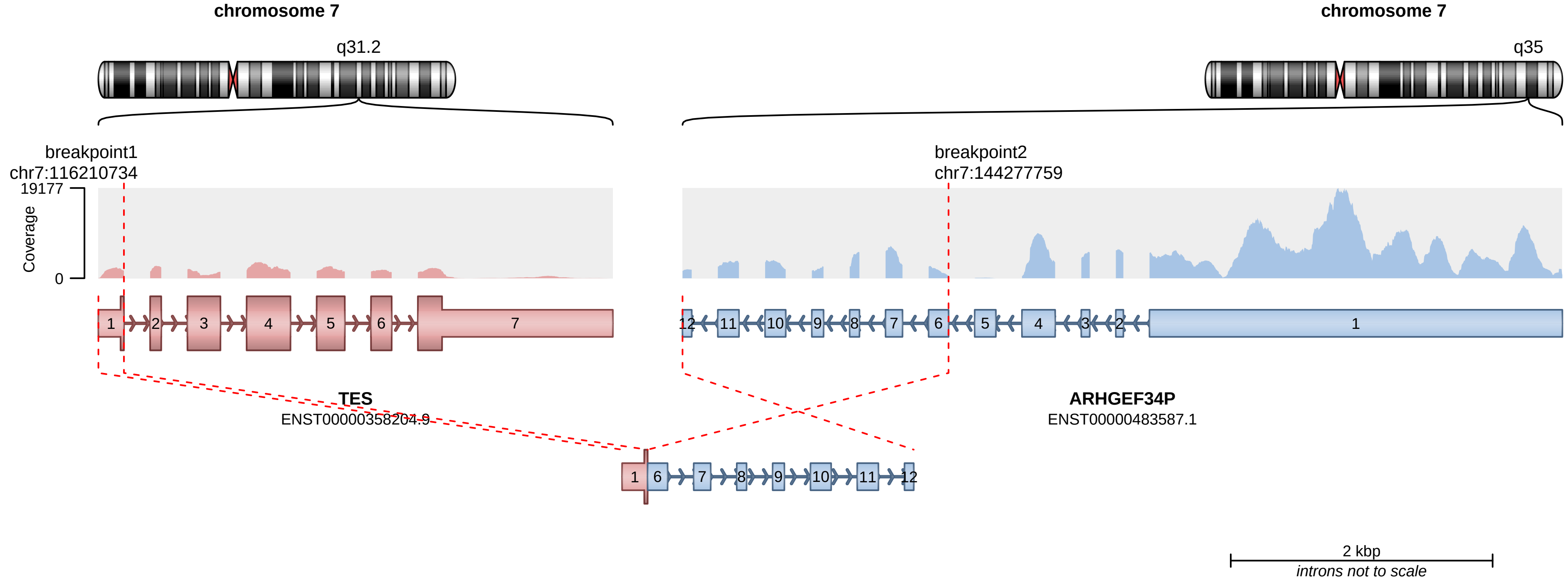


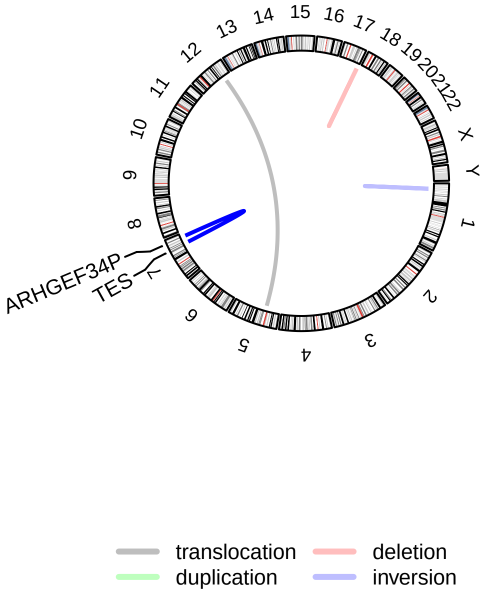
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 1

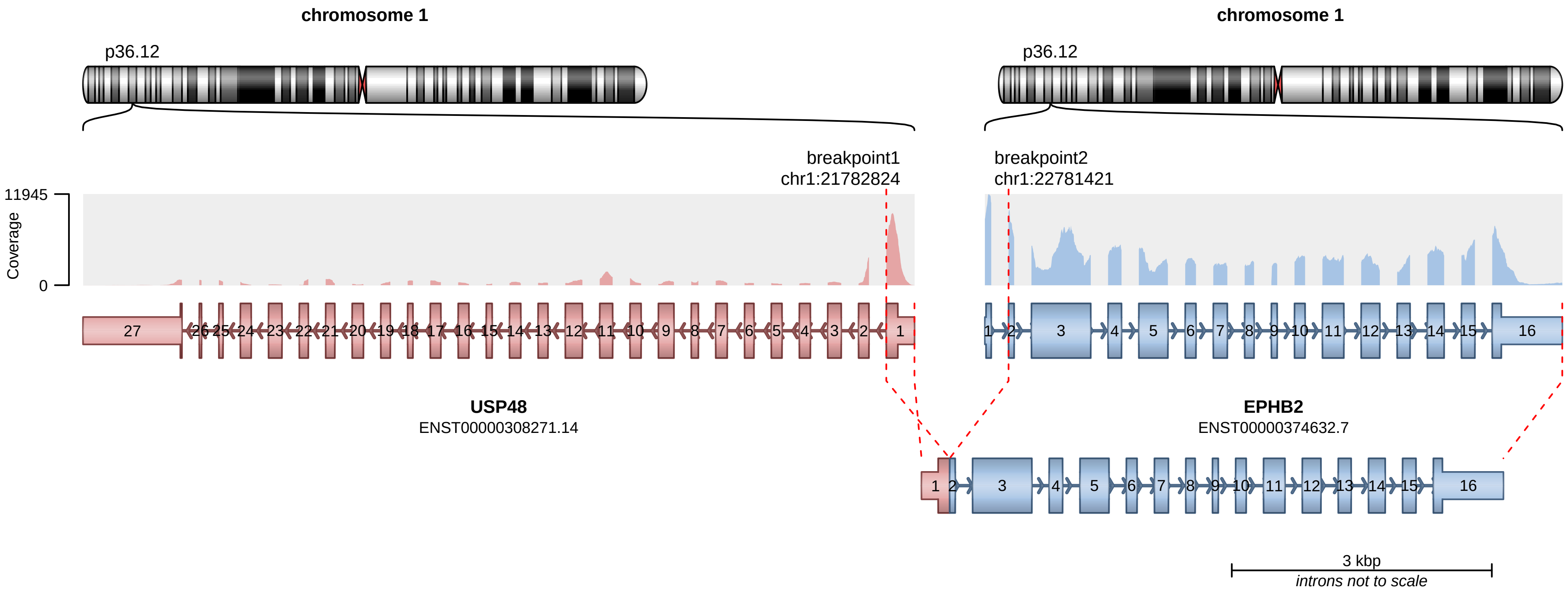


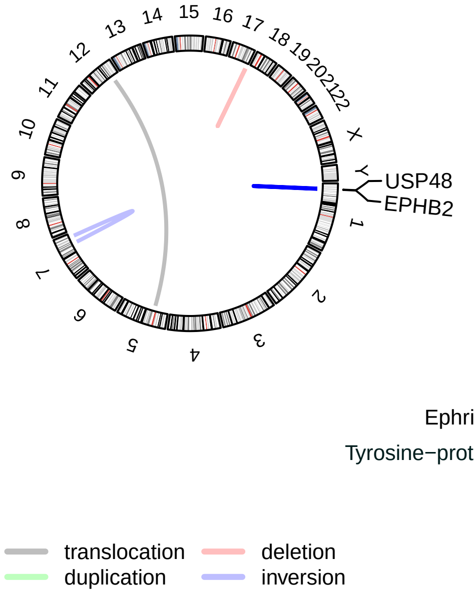


No protein domains retained in fusion.

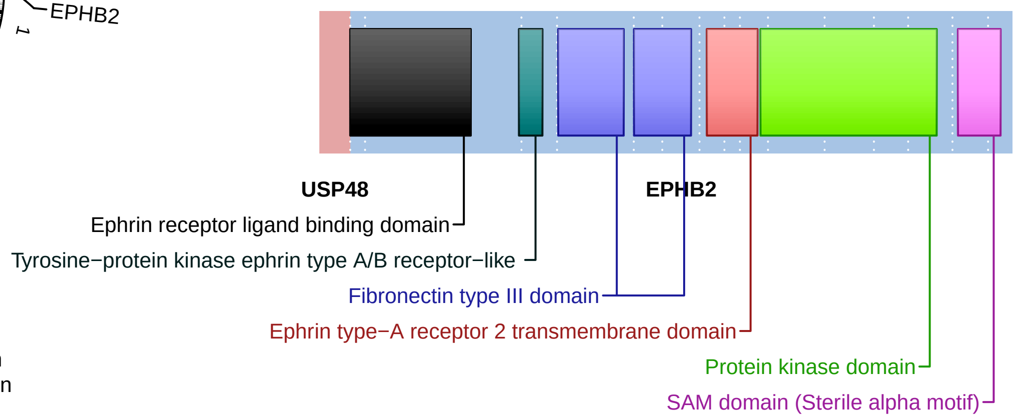
SUPPORTING READ COUNT

Split reads = 85
Discordant mates = 1



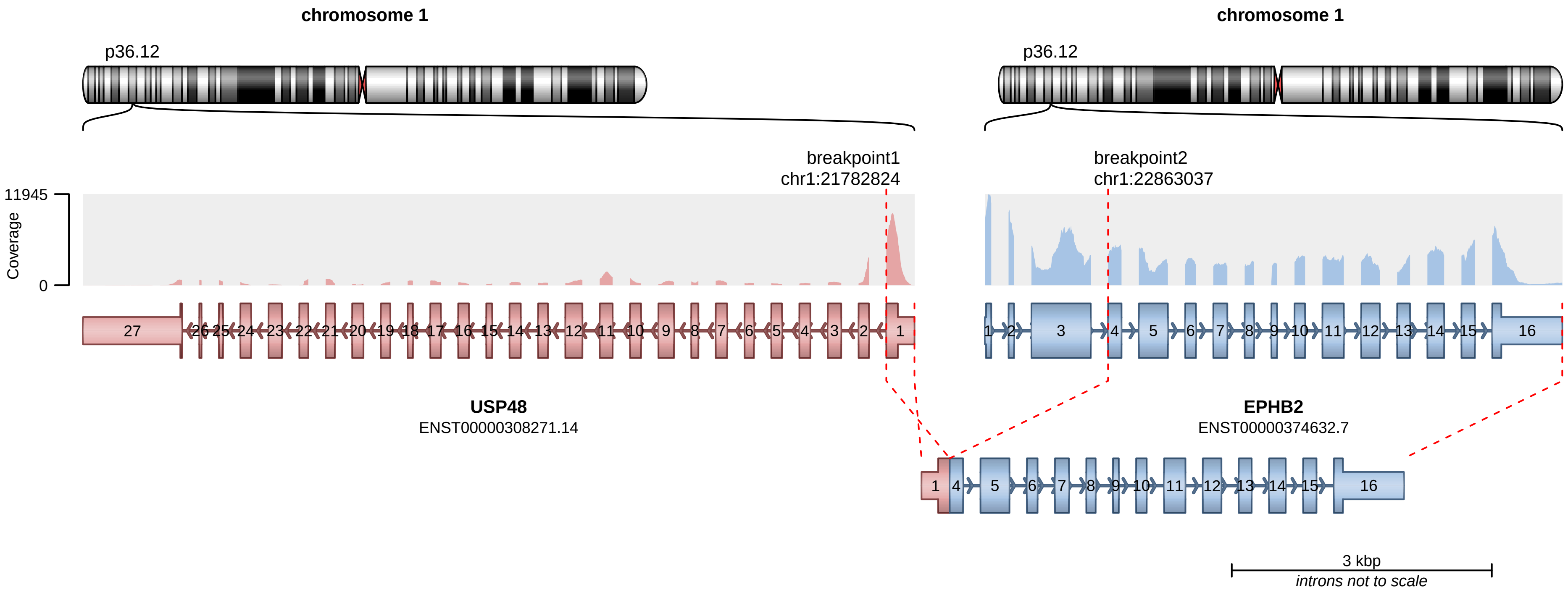


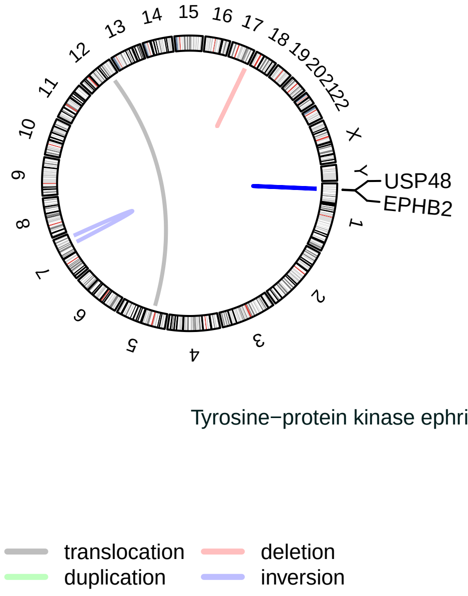
RETAINED PROTEIN DOMAINS
reading frame unclear

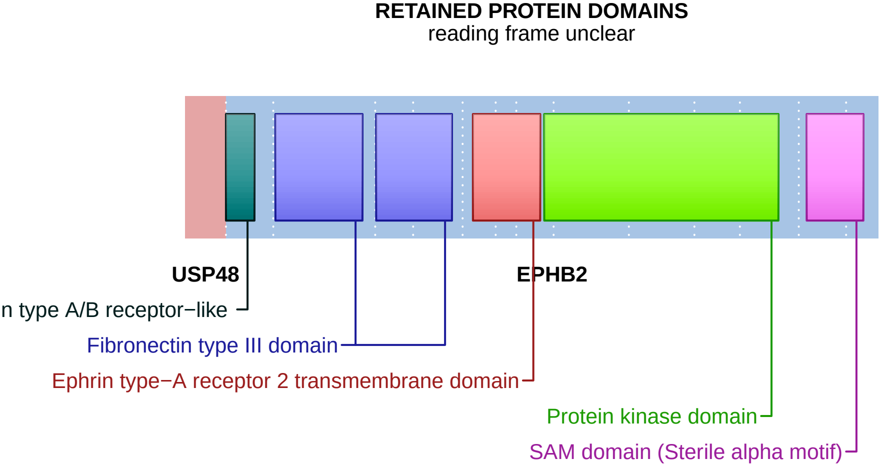


SUPPORTING READ COUNT

Split reads = 31
Discordant mates = 0

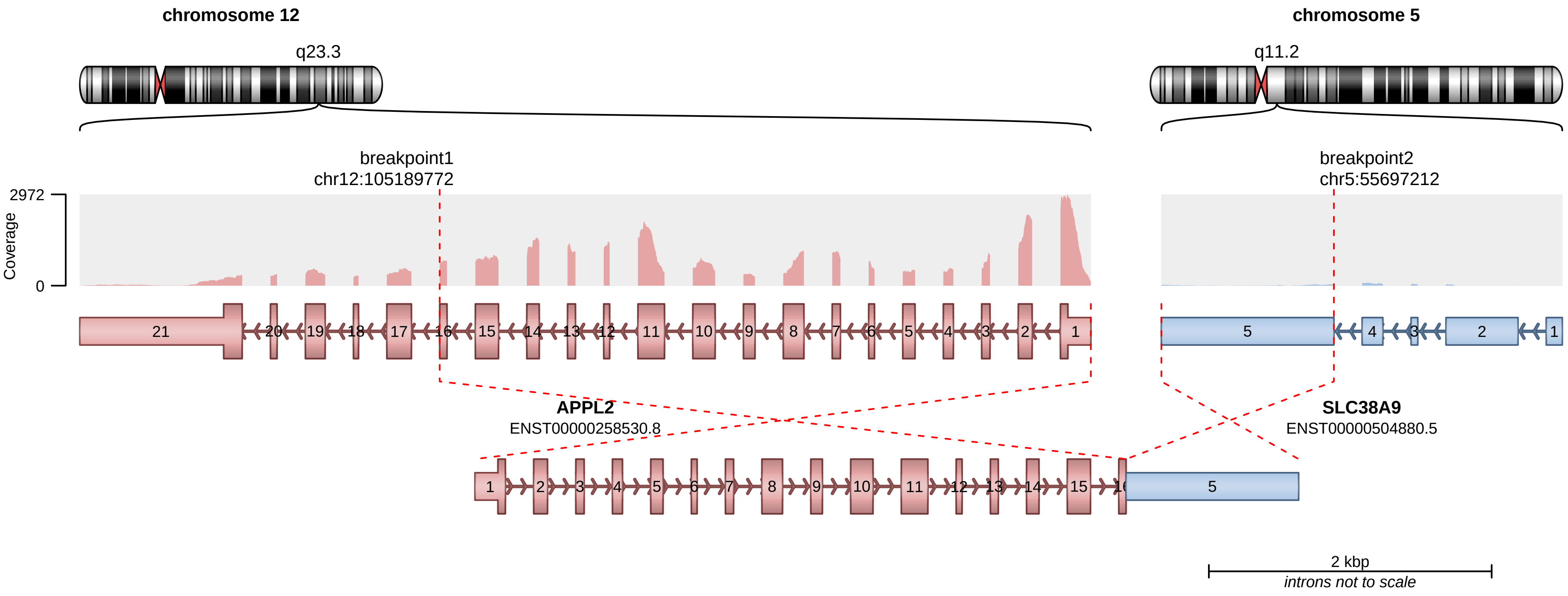


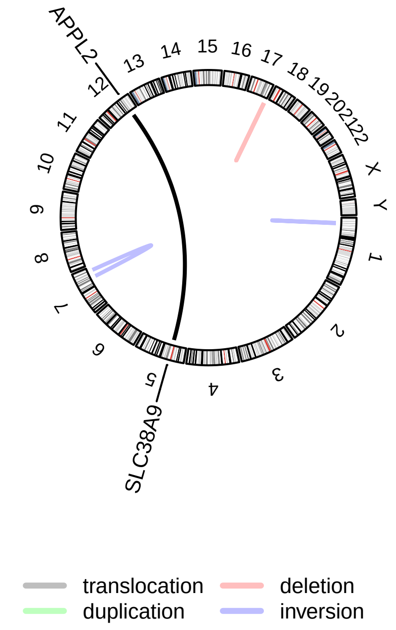




SUPPORTING READ COUNT

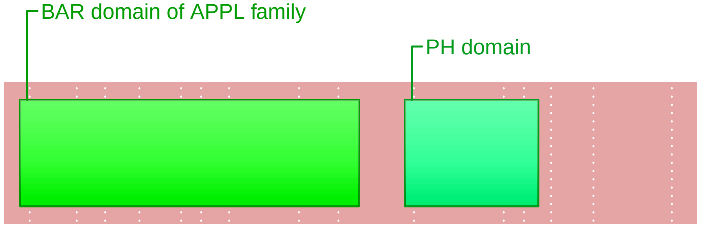
Split reads = 4
Discordant mates = 0





RETAINED PROTEIN DOMAINS

reading frame unclear



APPL2

SUPPORTING READ COUNT

Split reads = 11
Discordant mates = 0