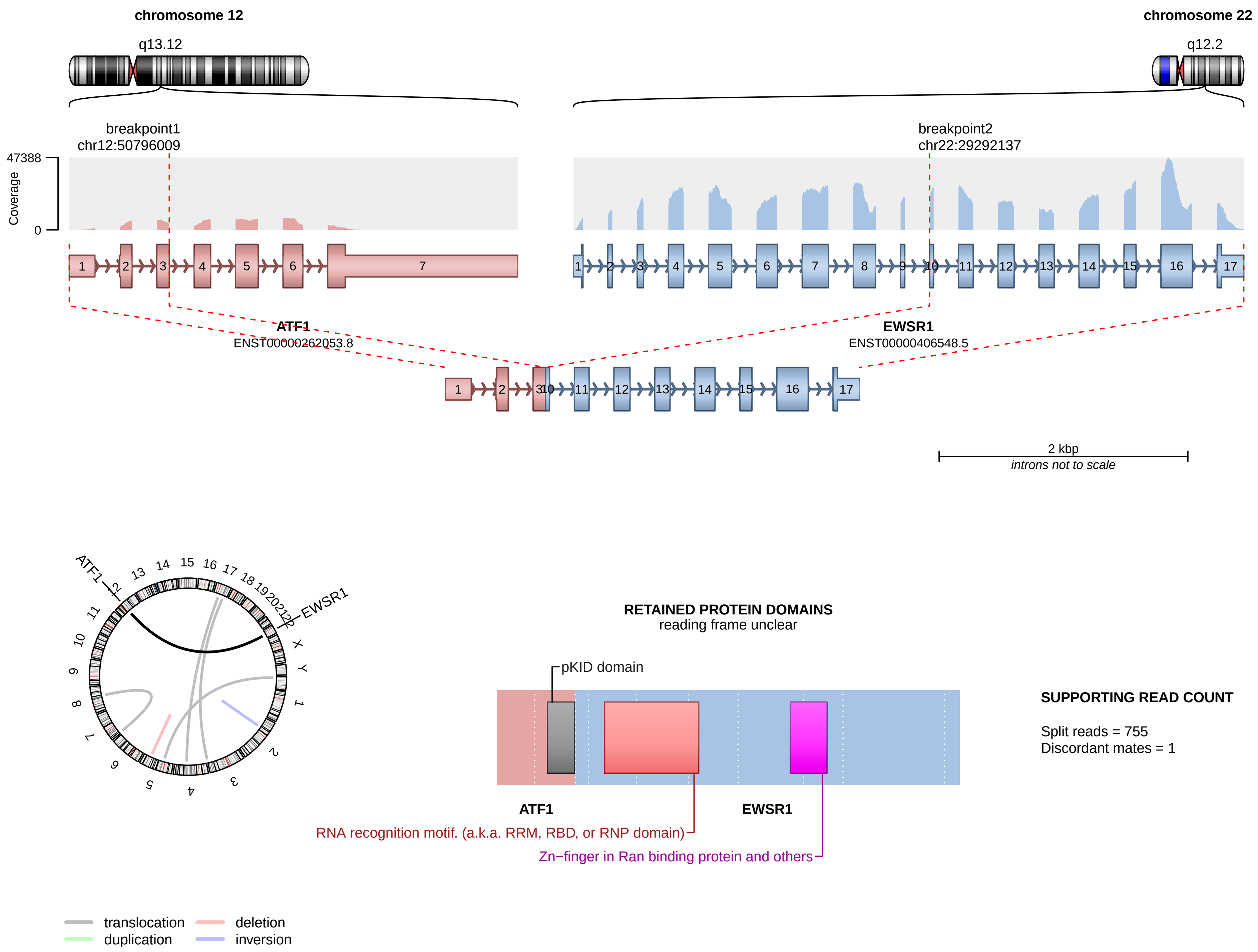
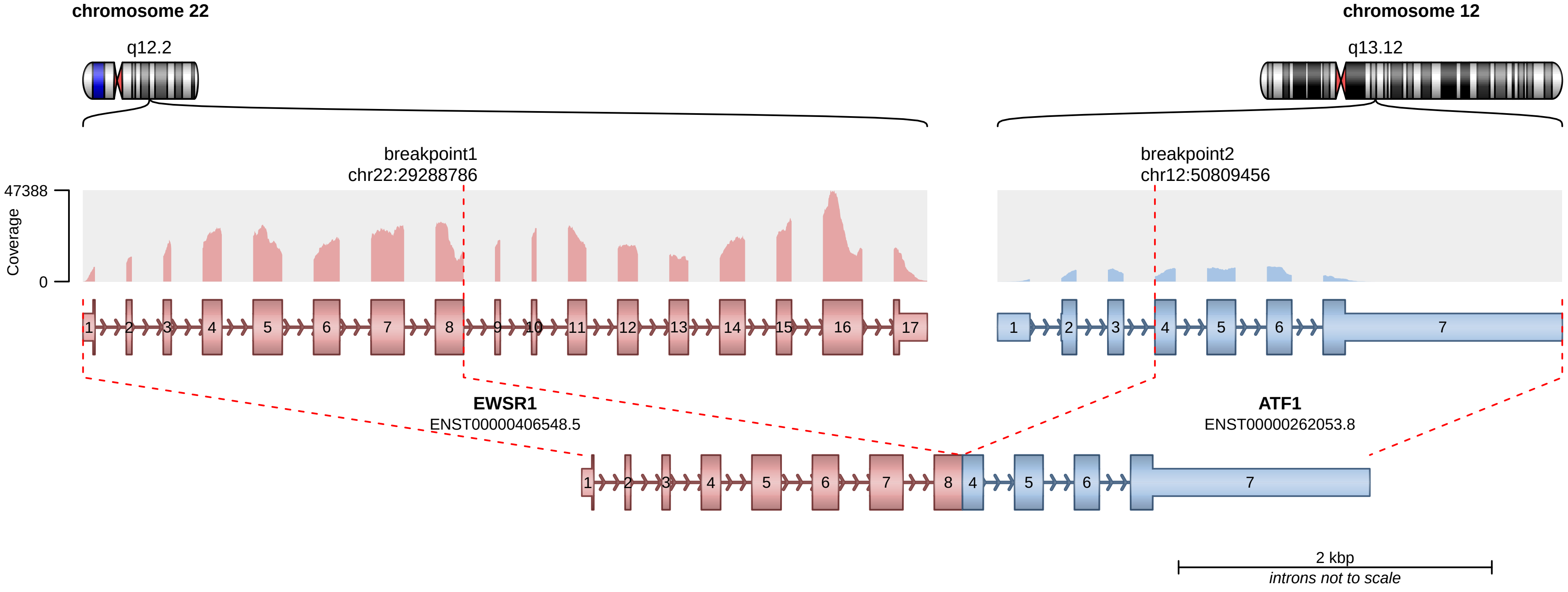
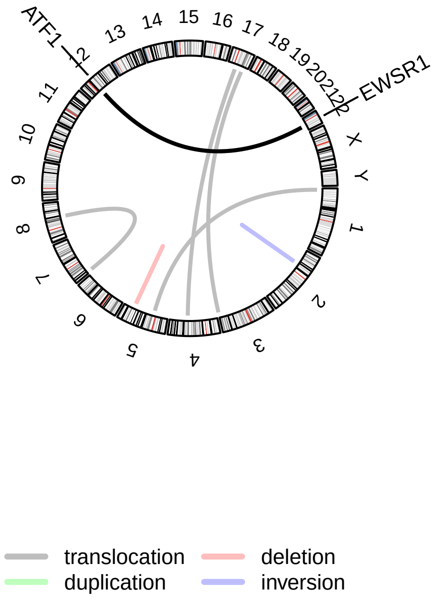
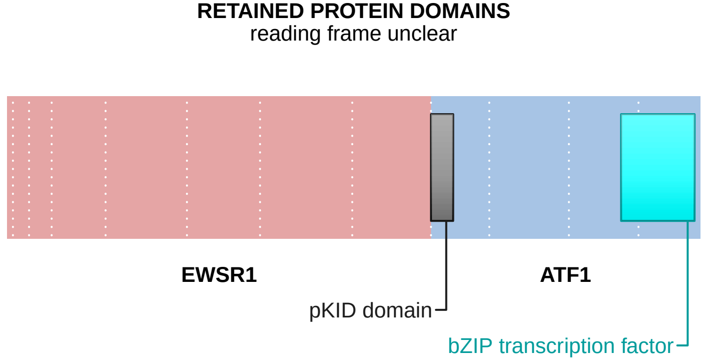


SUPPORTING READ COUNT

Split reads = 755
Discordant mates = 1

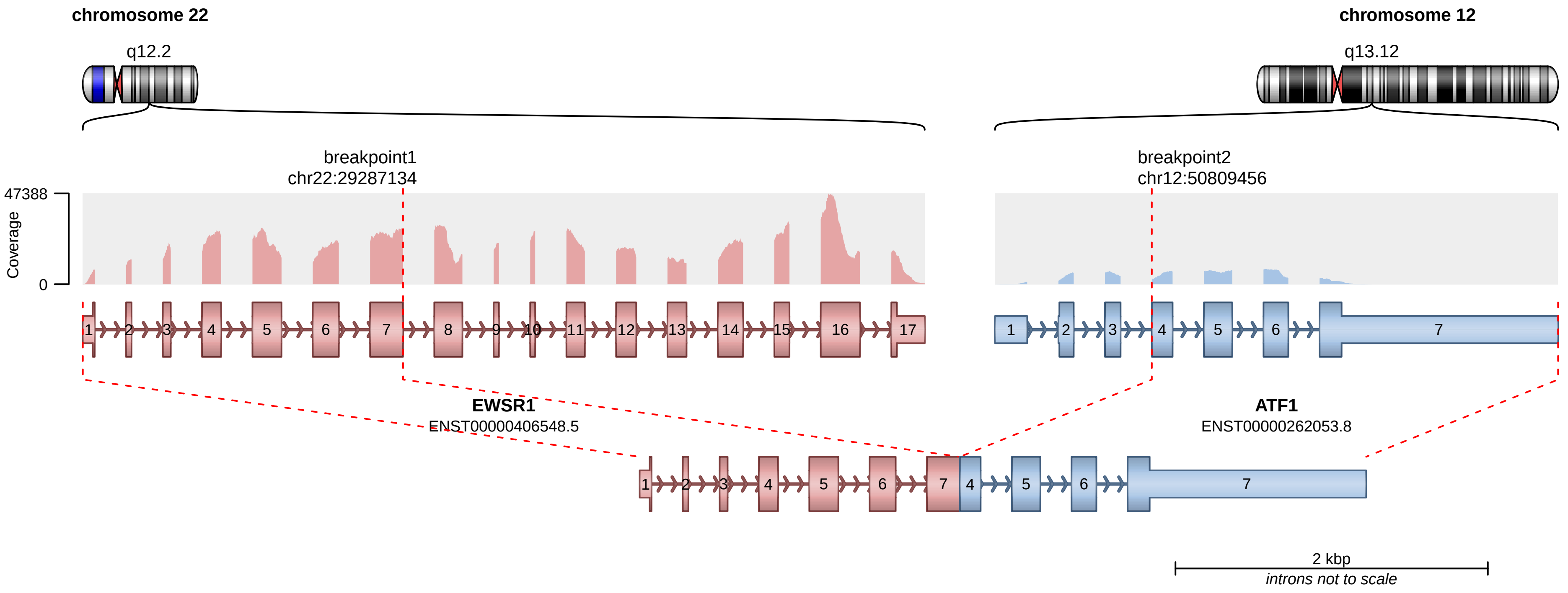


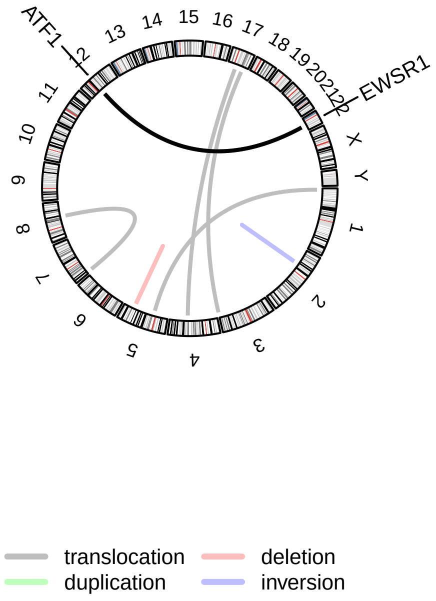


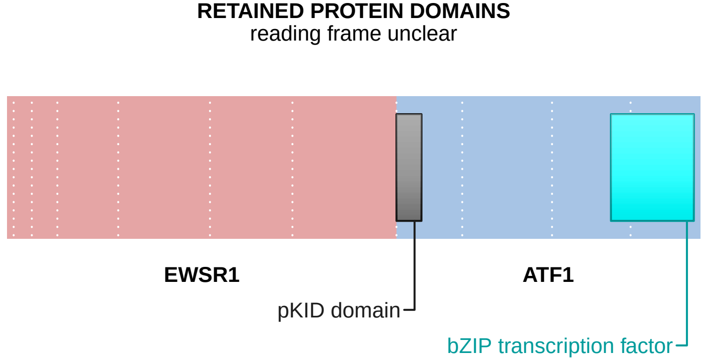


SUPPORTING READ COUNT

Split reads = 463
Discordant mates = 0

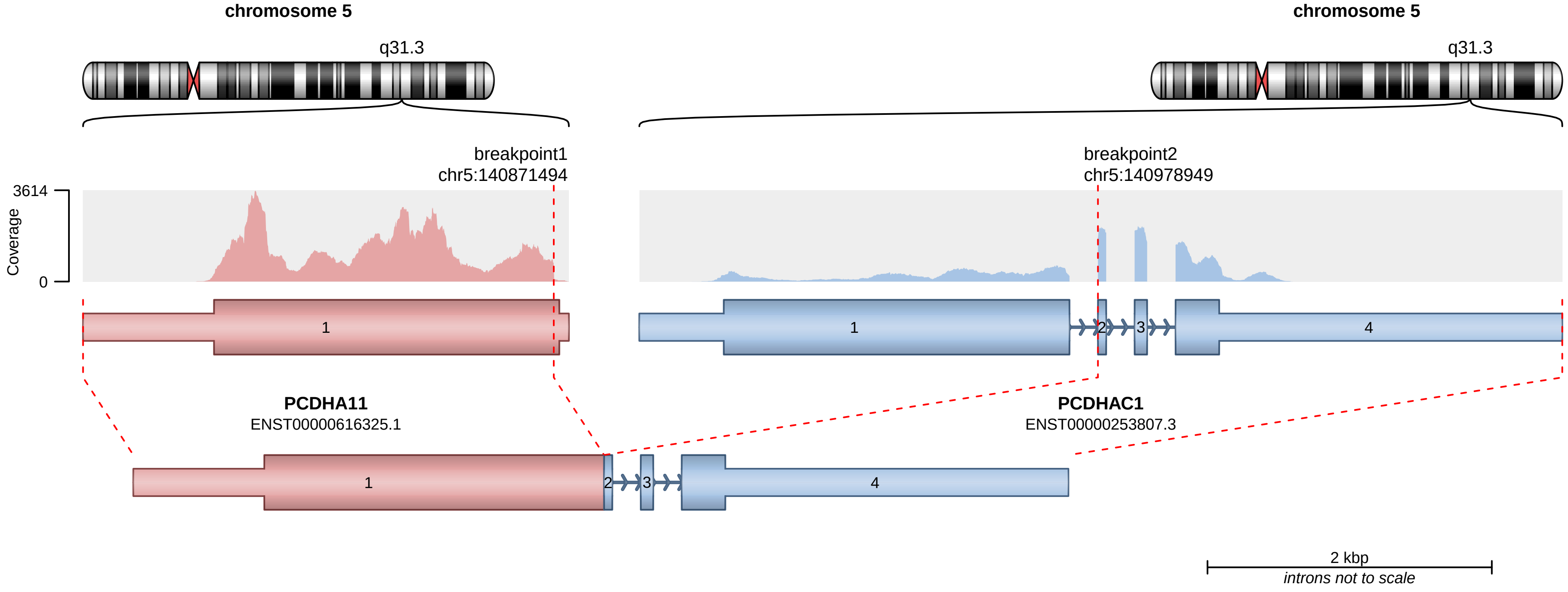


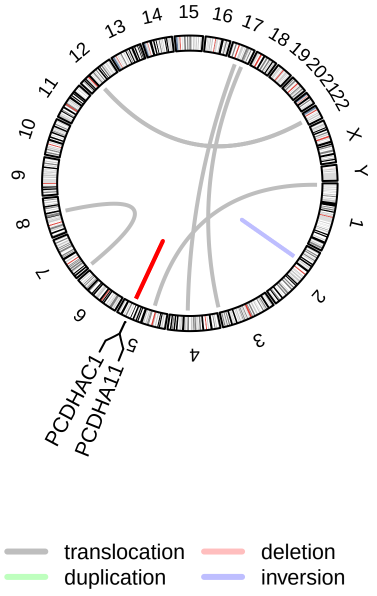


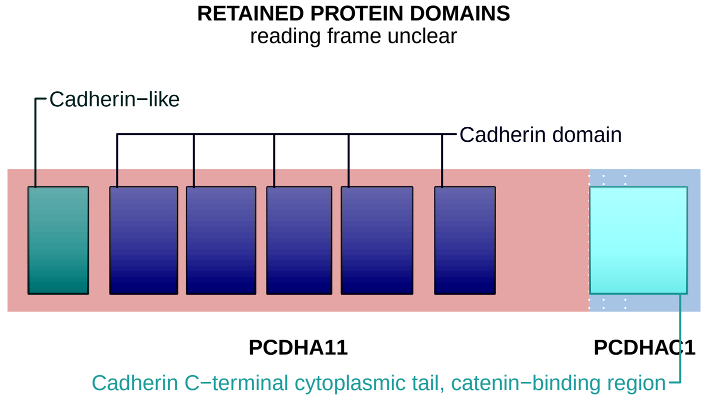


SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

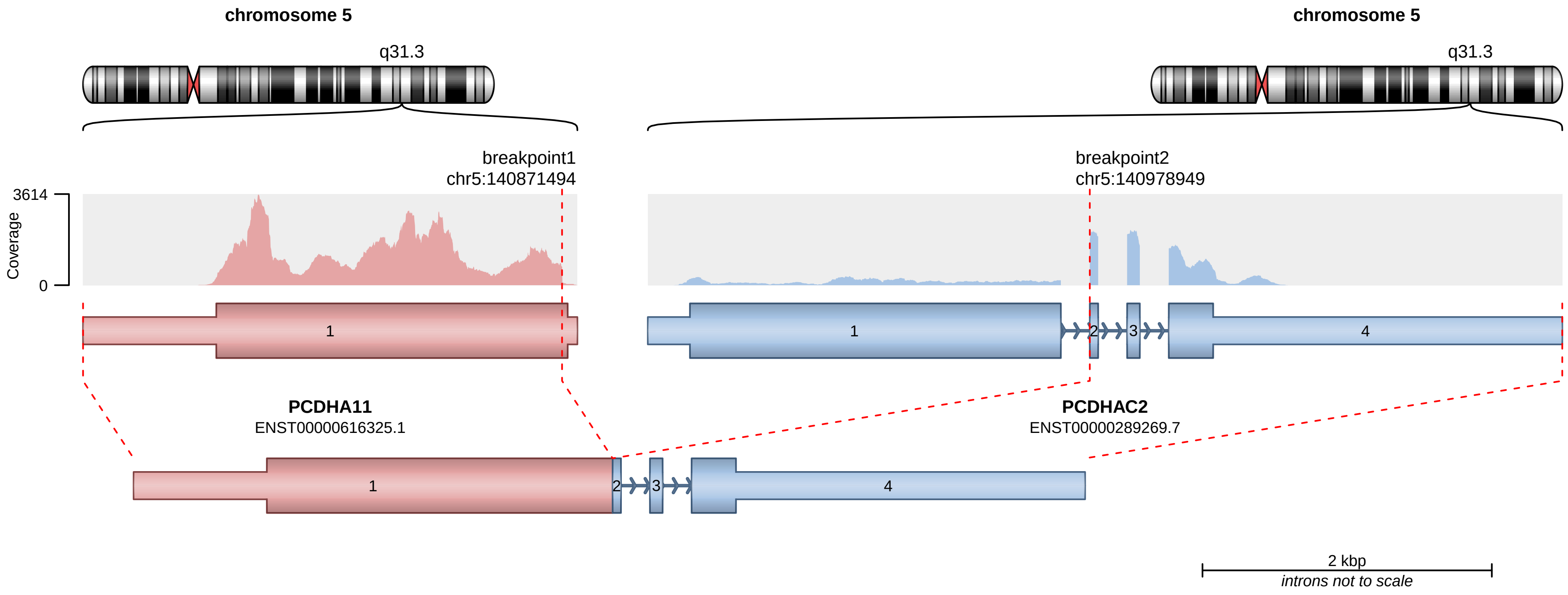


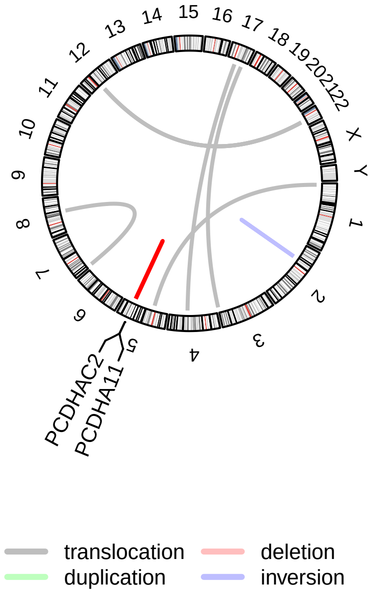




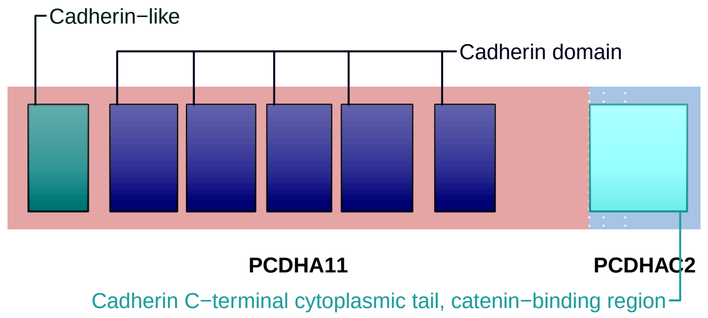
SUPPORTING READ COUNT

Split reads = 187
Discordant mates = 0



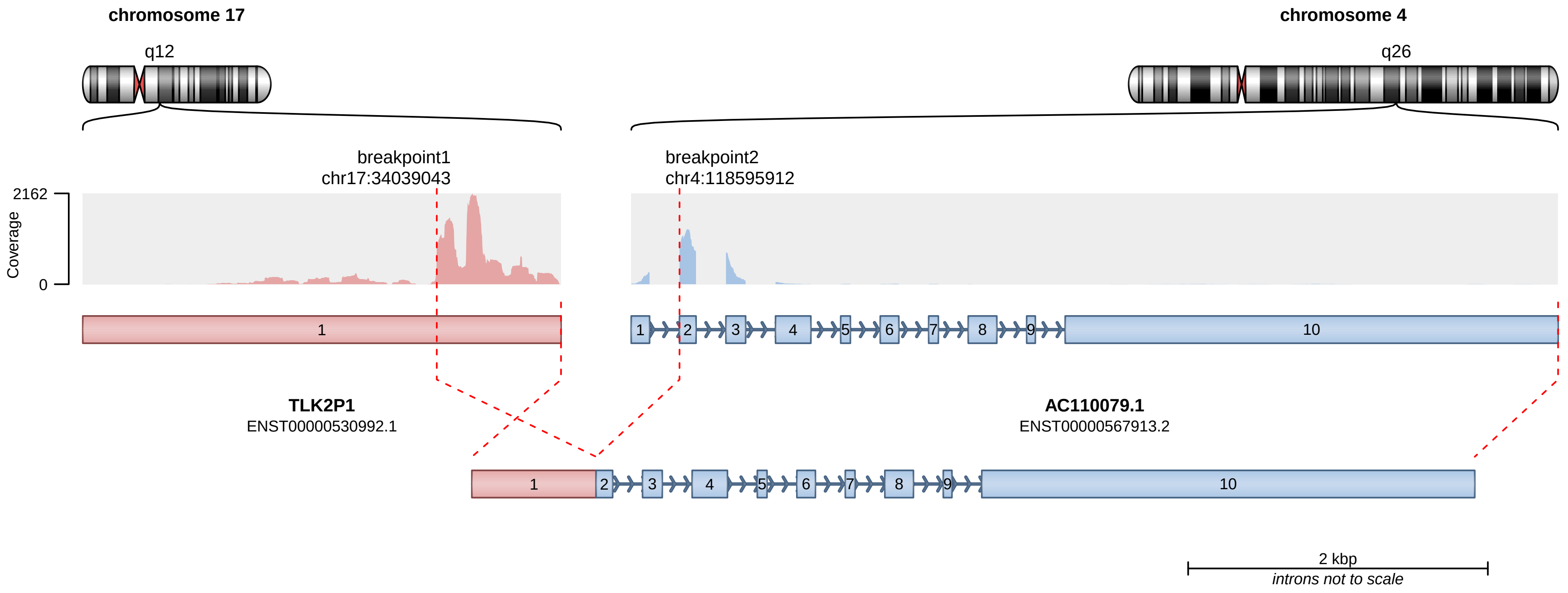


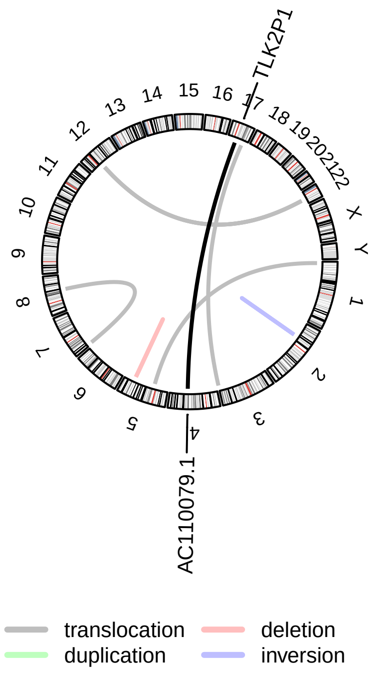
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 187
Discordant mates = 0

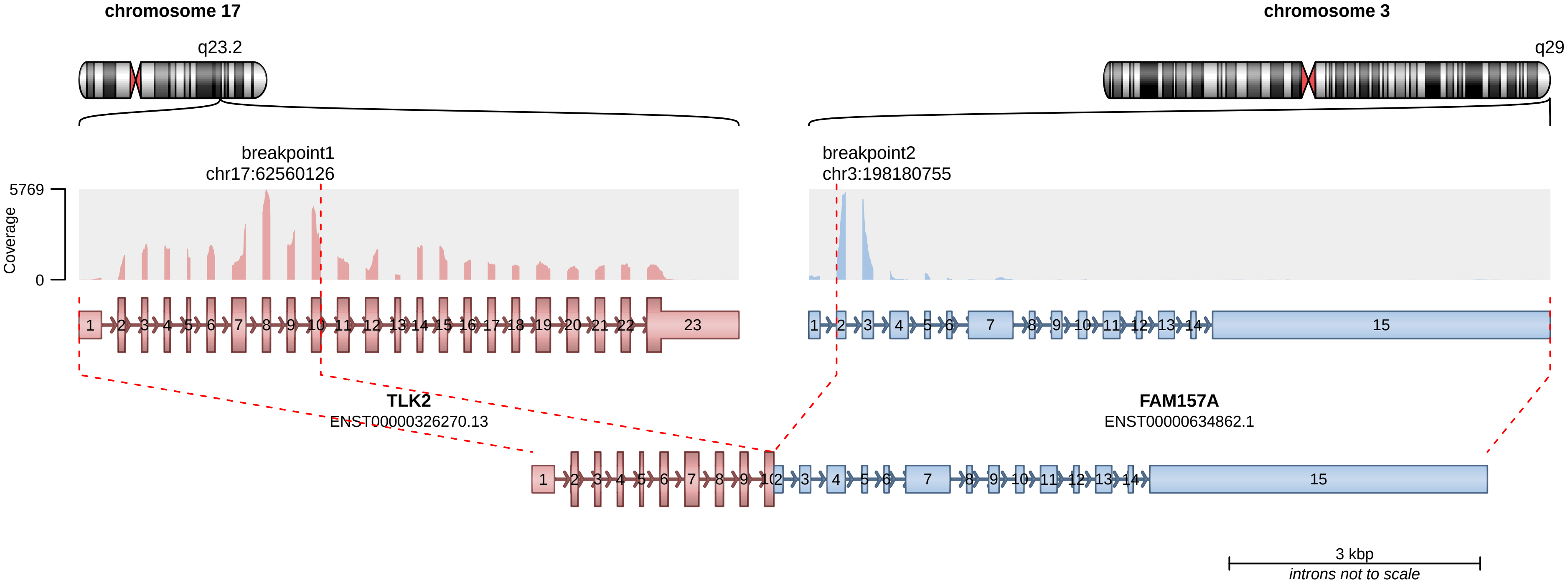


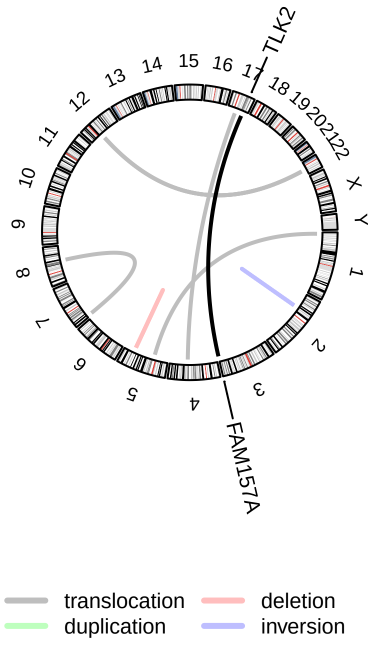


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 141
Discordant mates = 0

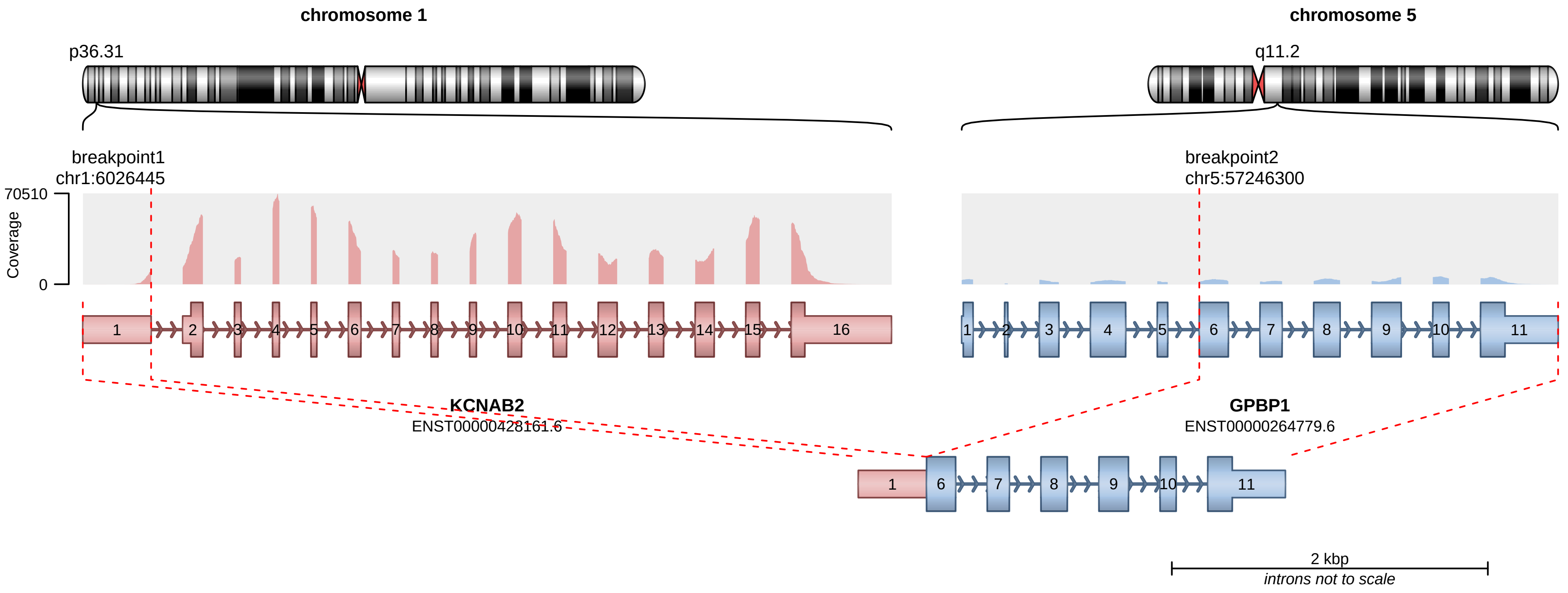


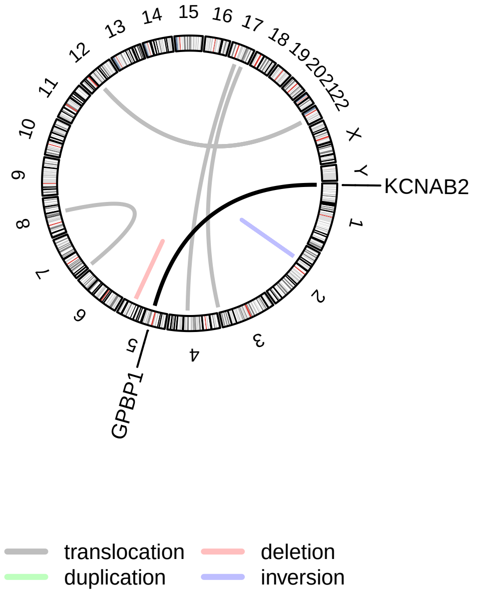


No protein domains retained in fusion.

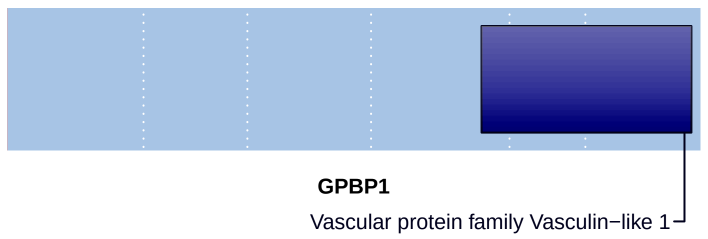
SUPPORTING READ COUNT

Split reads = 139
Discordant mates = 0



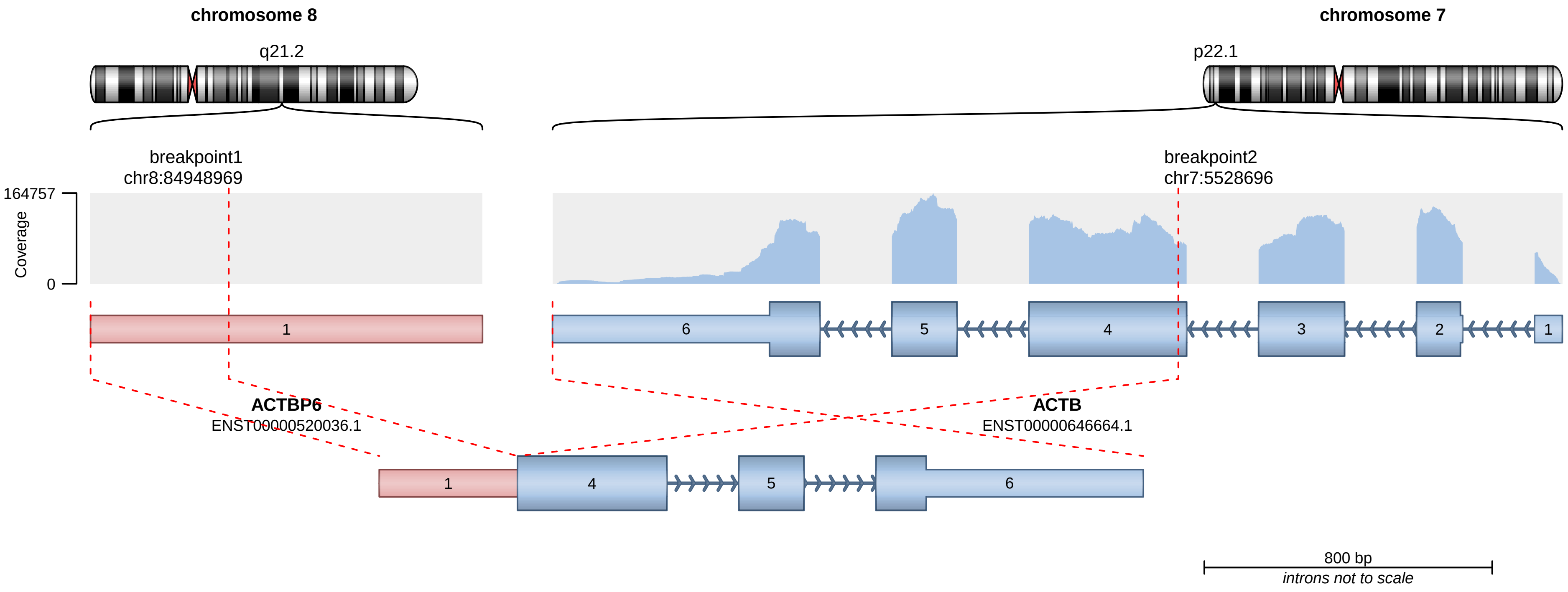


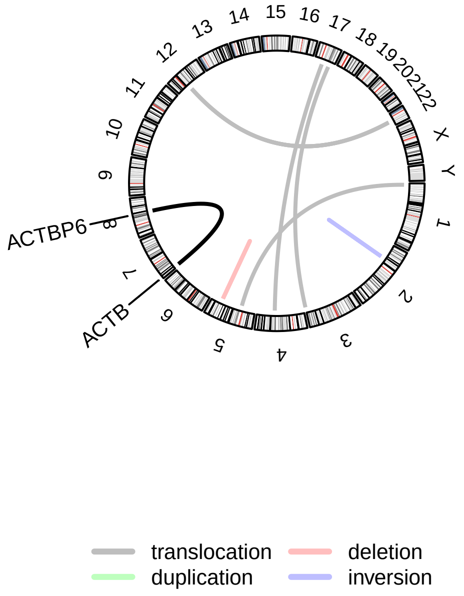
RETAINED PROTEIN DOMAINS
reading frame unclear

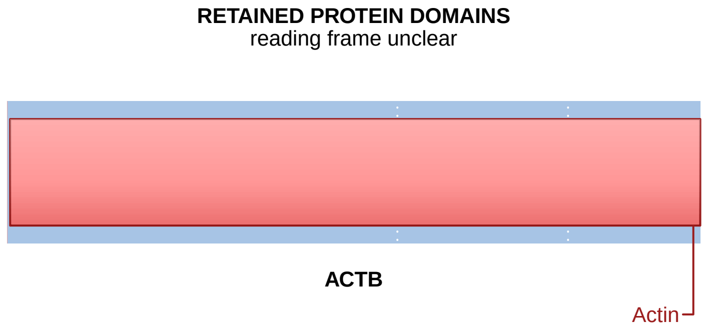


SUPPORTING READ COUNT

Split reads = 32
Discordant mates = 0

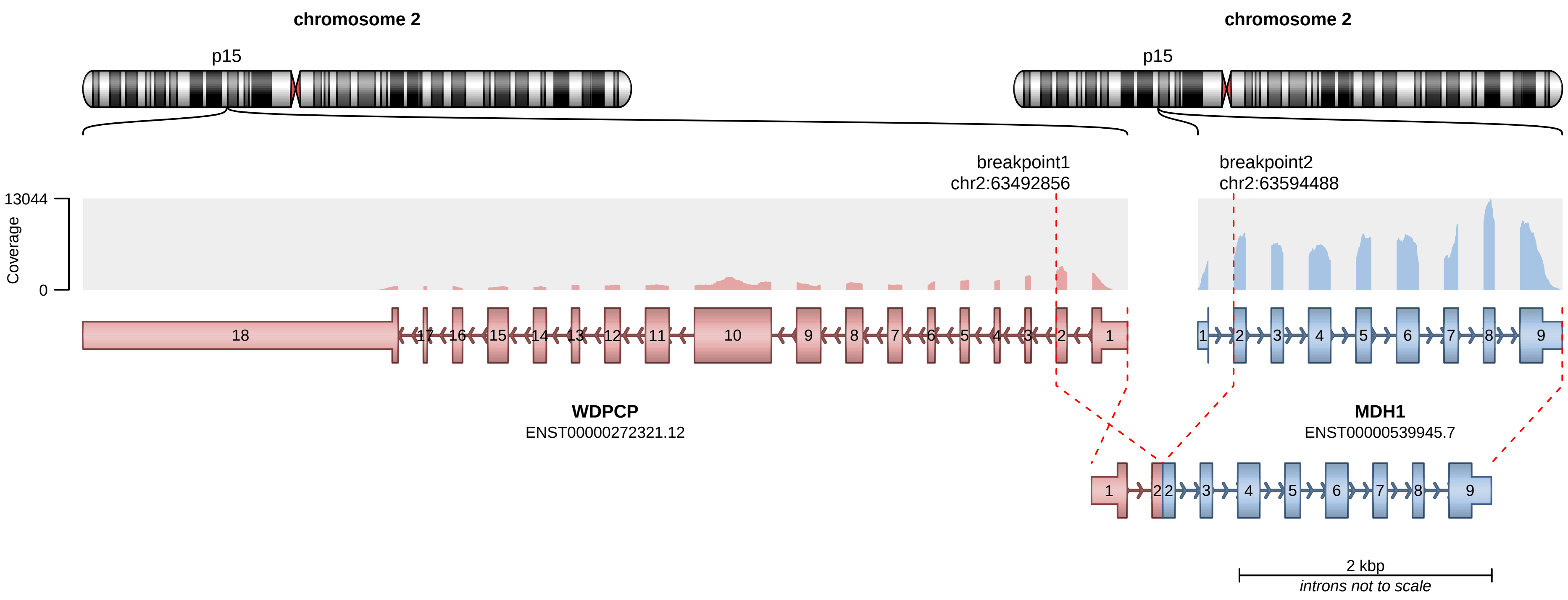


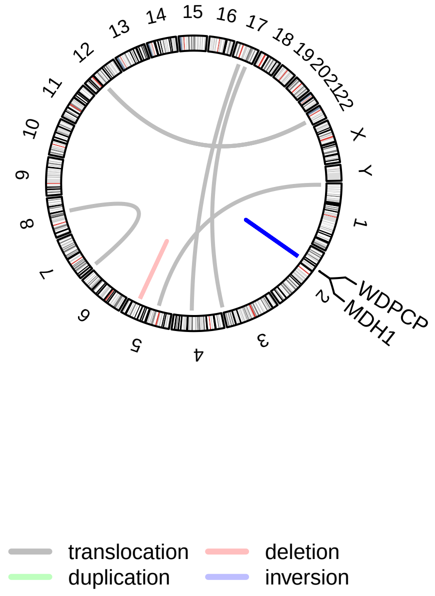




SUPPORTING READ COUNT

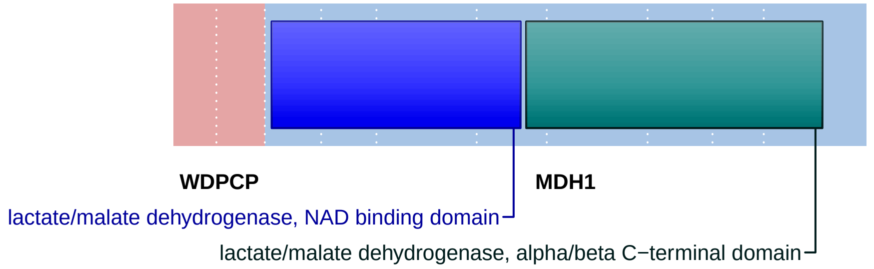
Split reads = 10
Discordant mates = 0





RETAINED PROTEIN DOMAINS

reading frame unclear



SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0