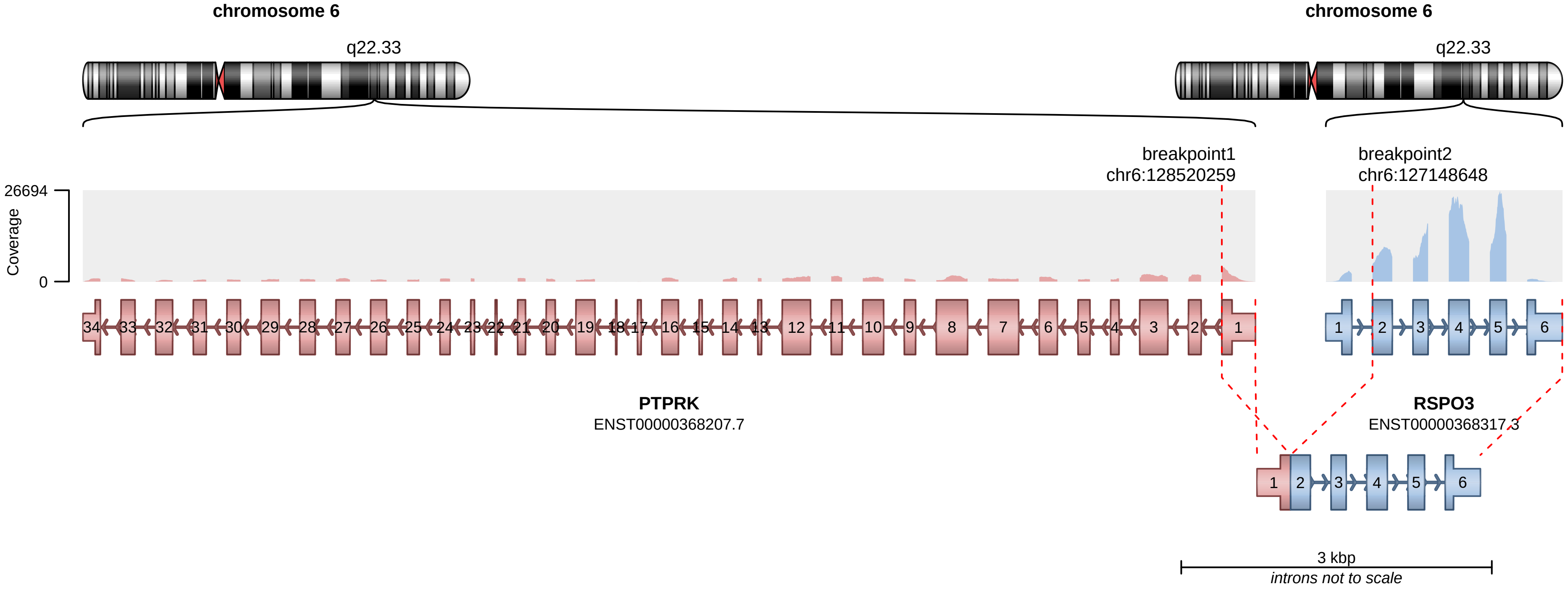
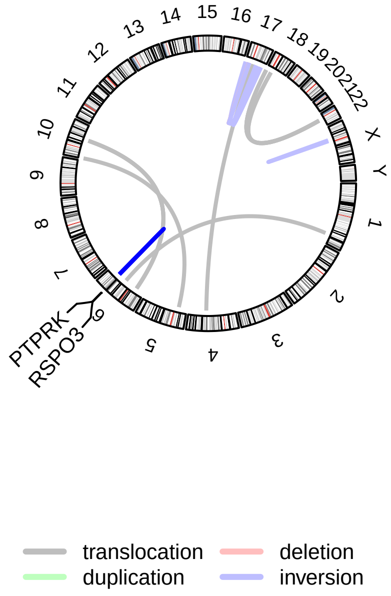
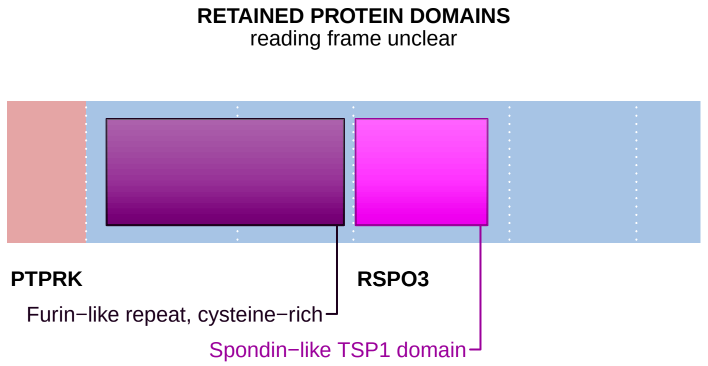
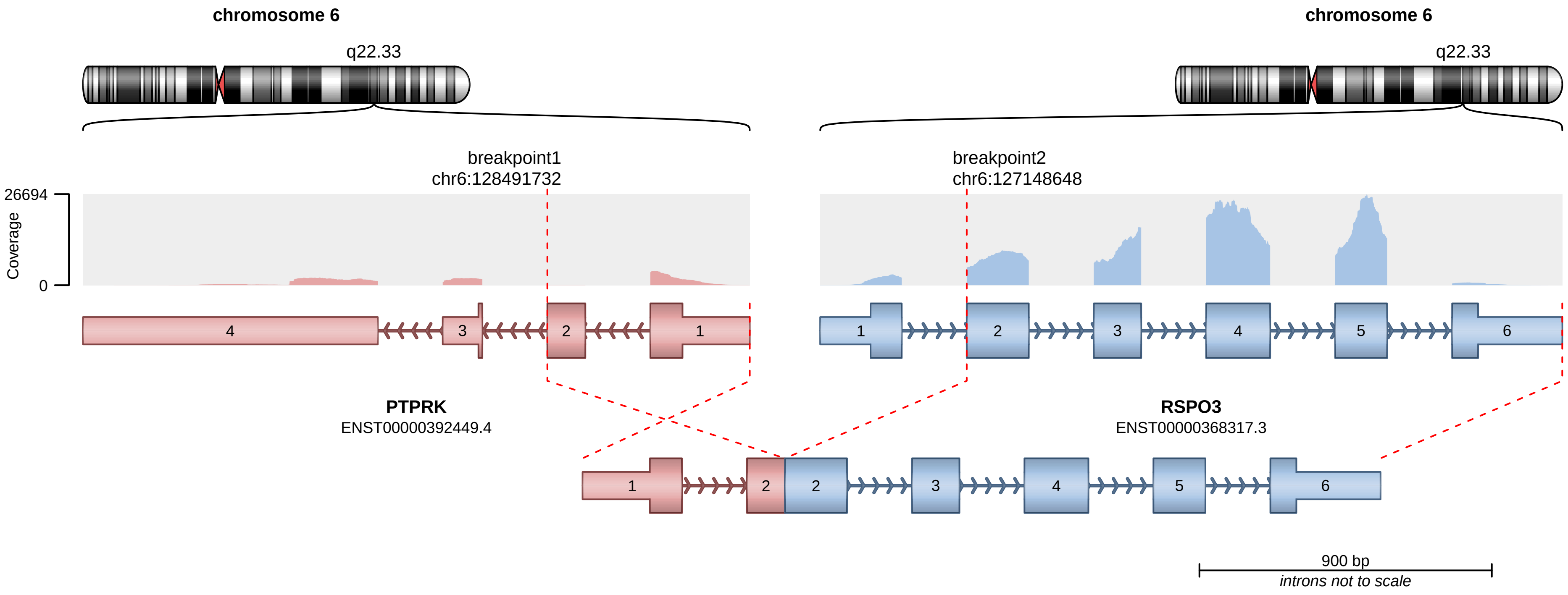


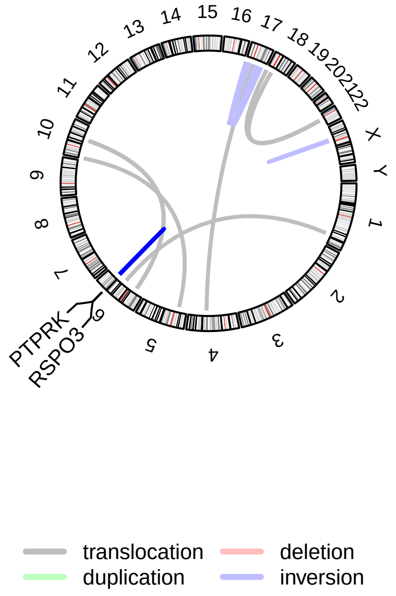


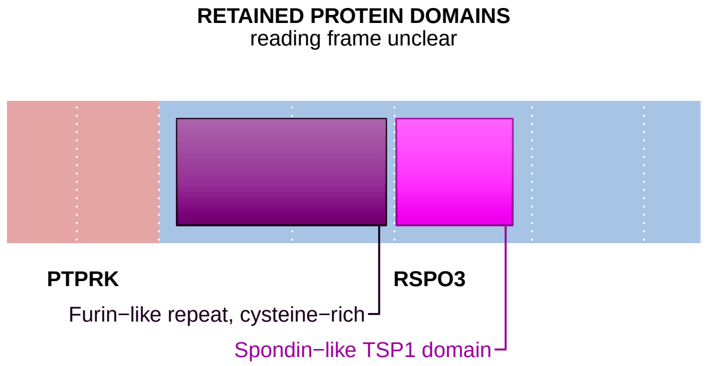


SUPPORTING READ COUNT

Split reads = 1241
Discordant mates = 1

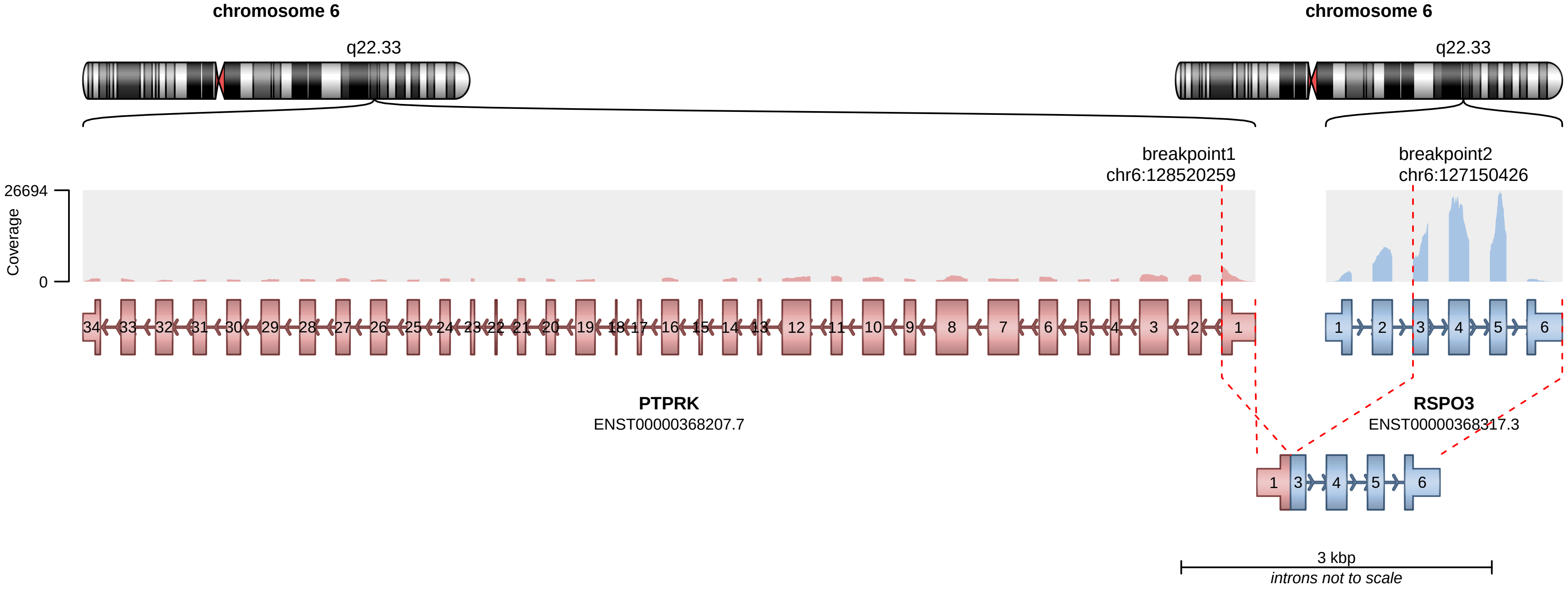


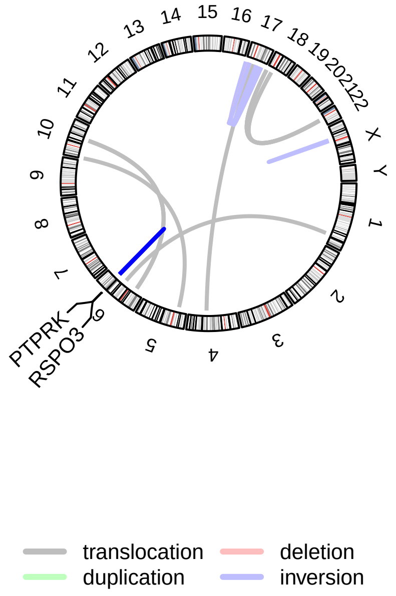


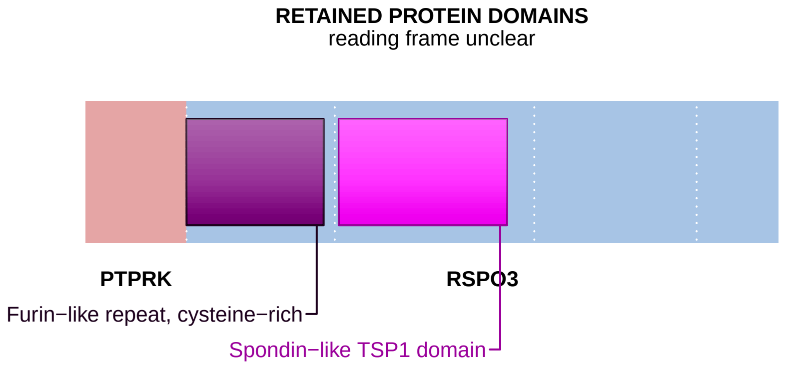


SUPPORTING READ COUNT

Split reads = 15
Discordant mates = 1

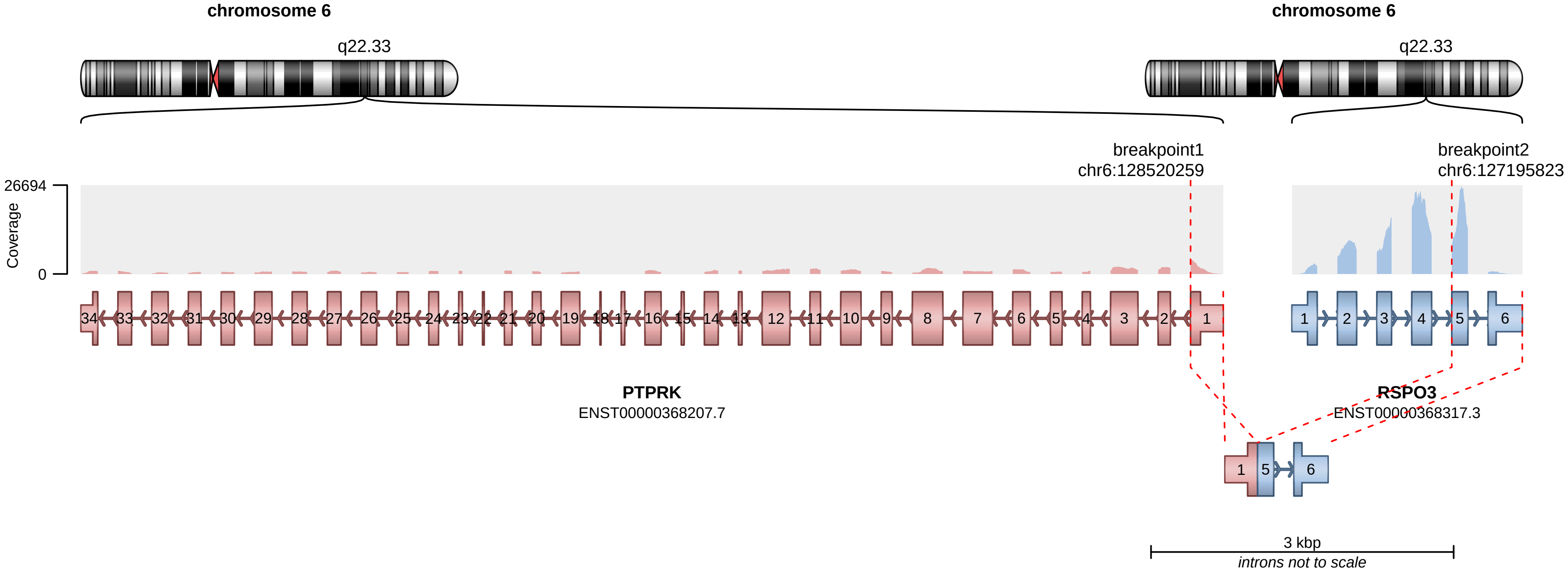


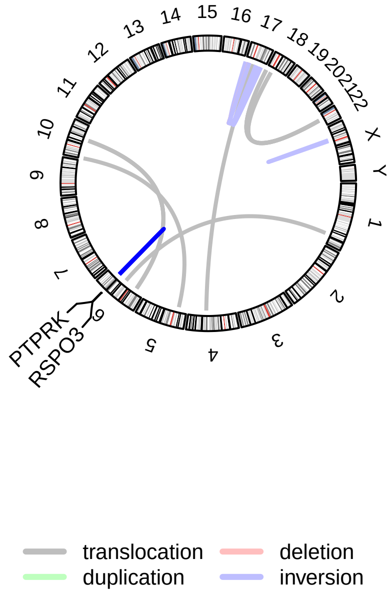




SUPPORTING READ COUNT

Split reads = 5
Discordant mates = 0

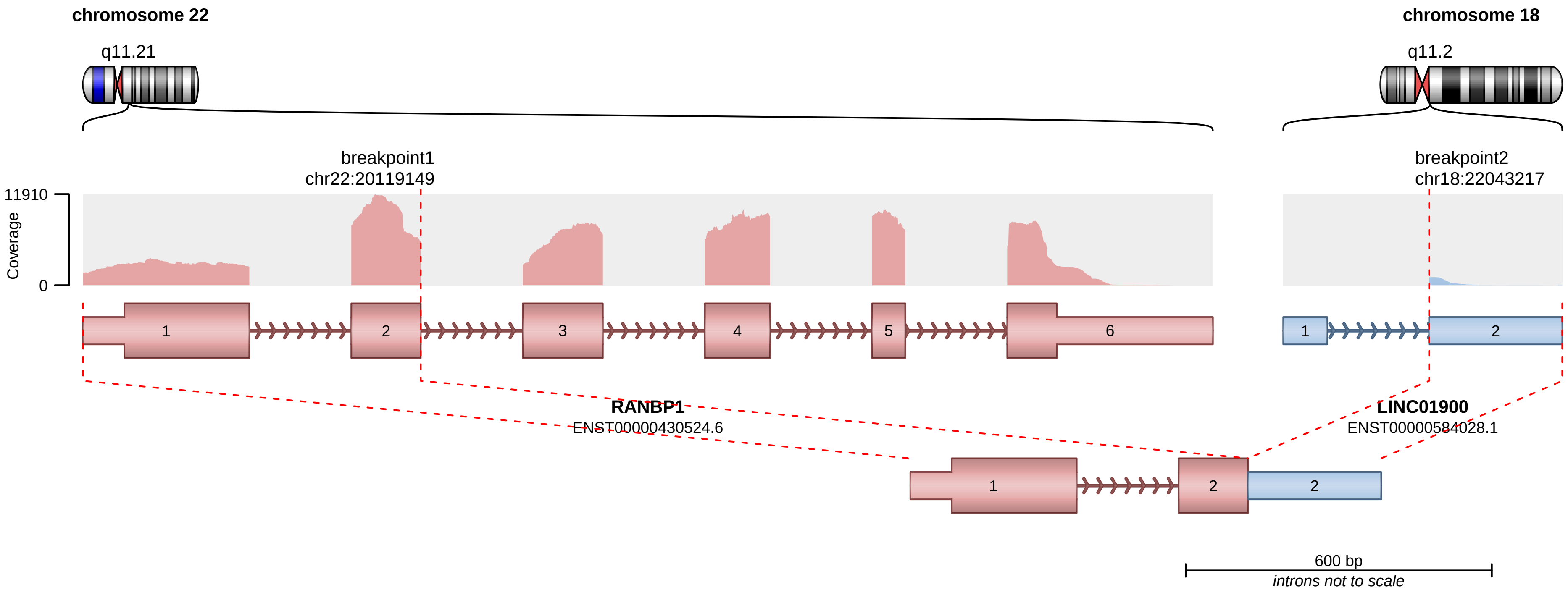


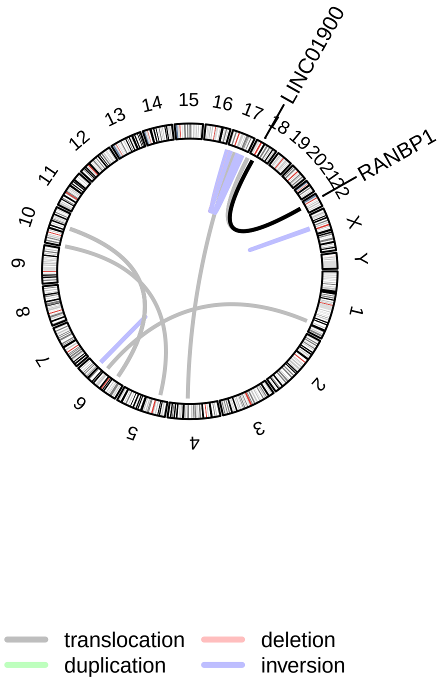


No protein domains retained in fusion.

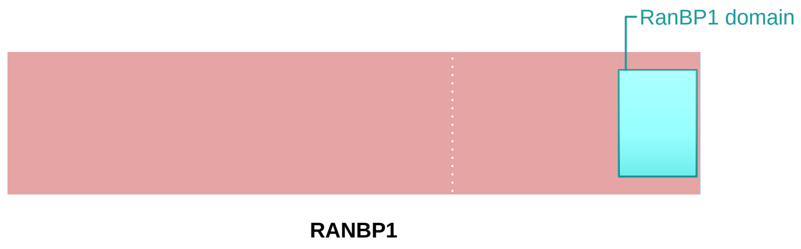
SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0



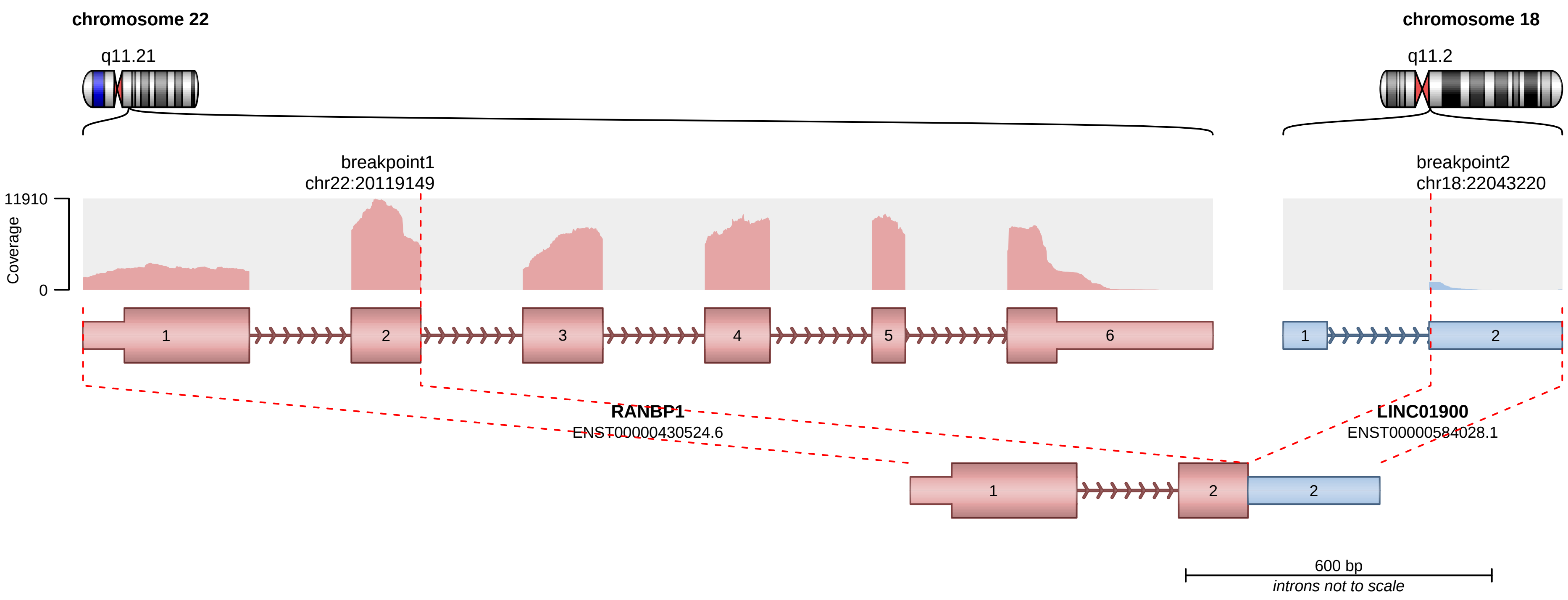


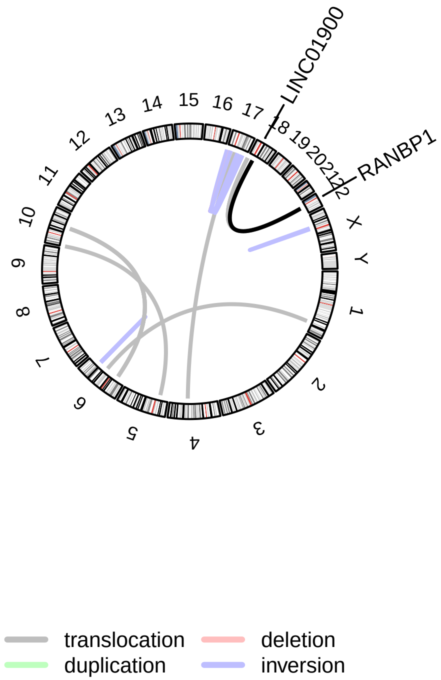
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 779
Discordant mates = 0



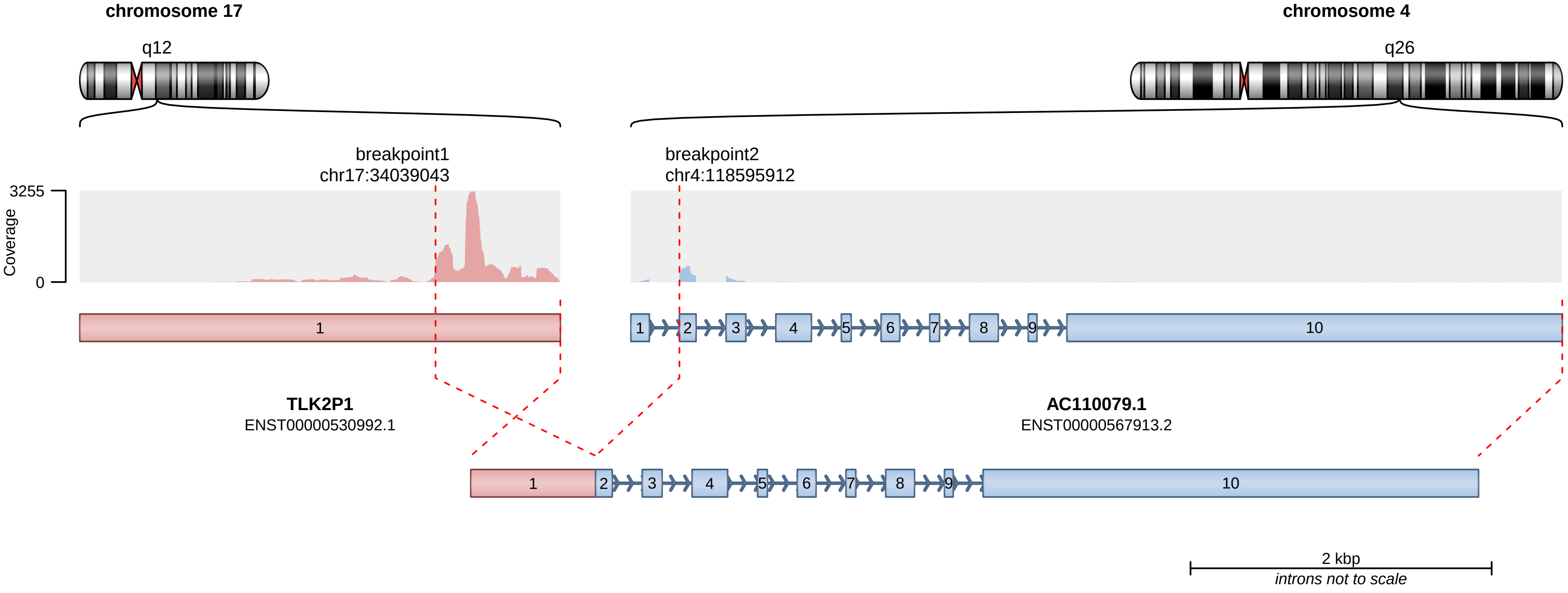


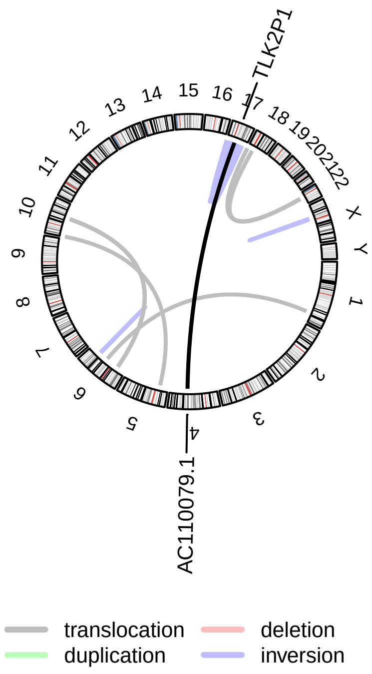
RETAINED PROTEIN DOMAINS
reading frame unclear

RANBP1

SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 0

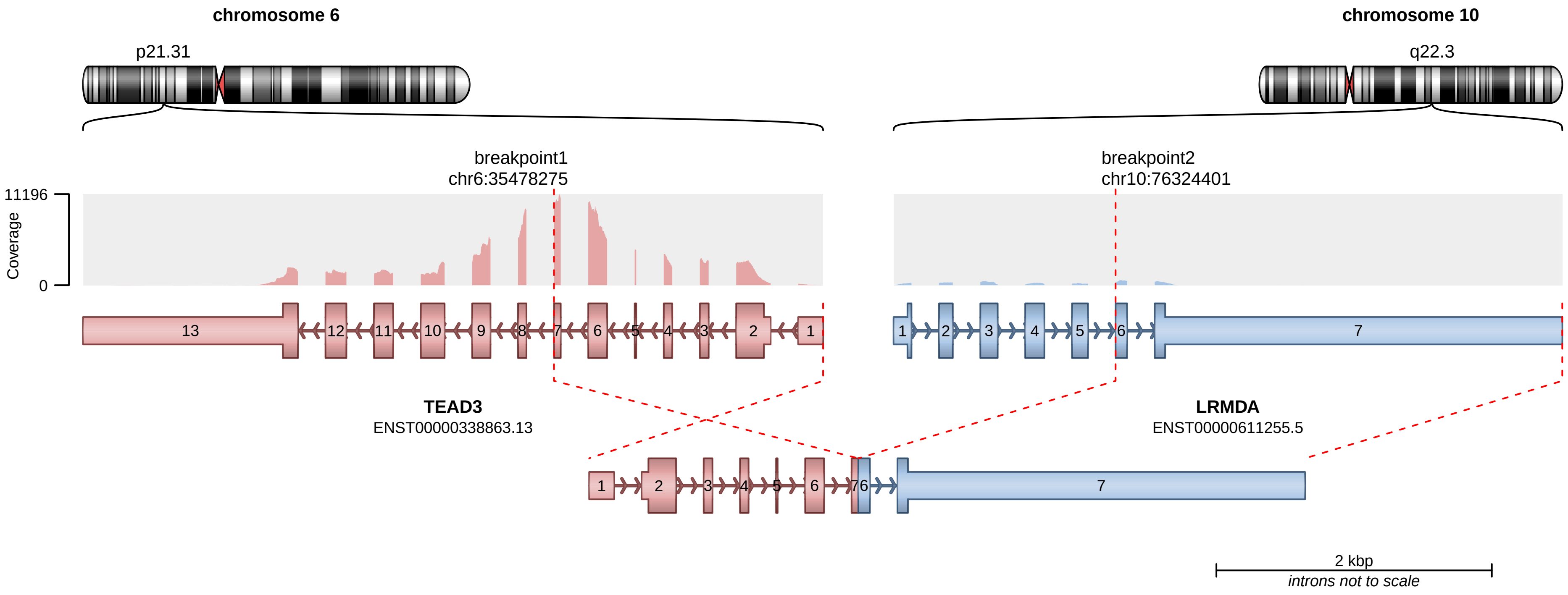


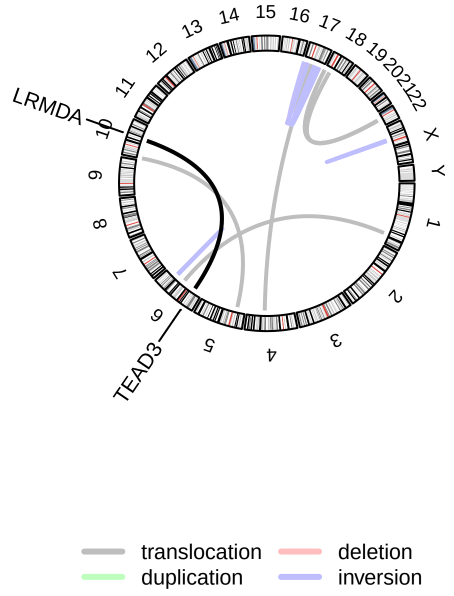


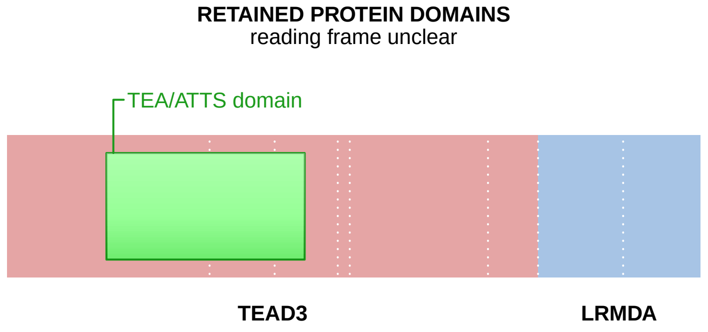
Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 83
Discordant mates = 0

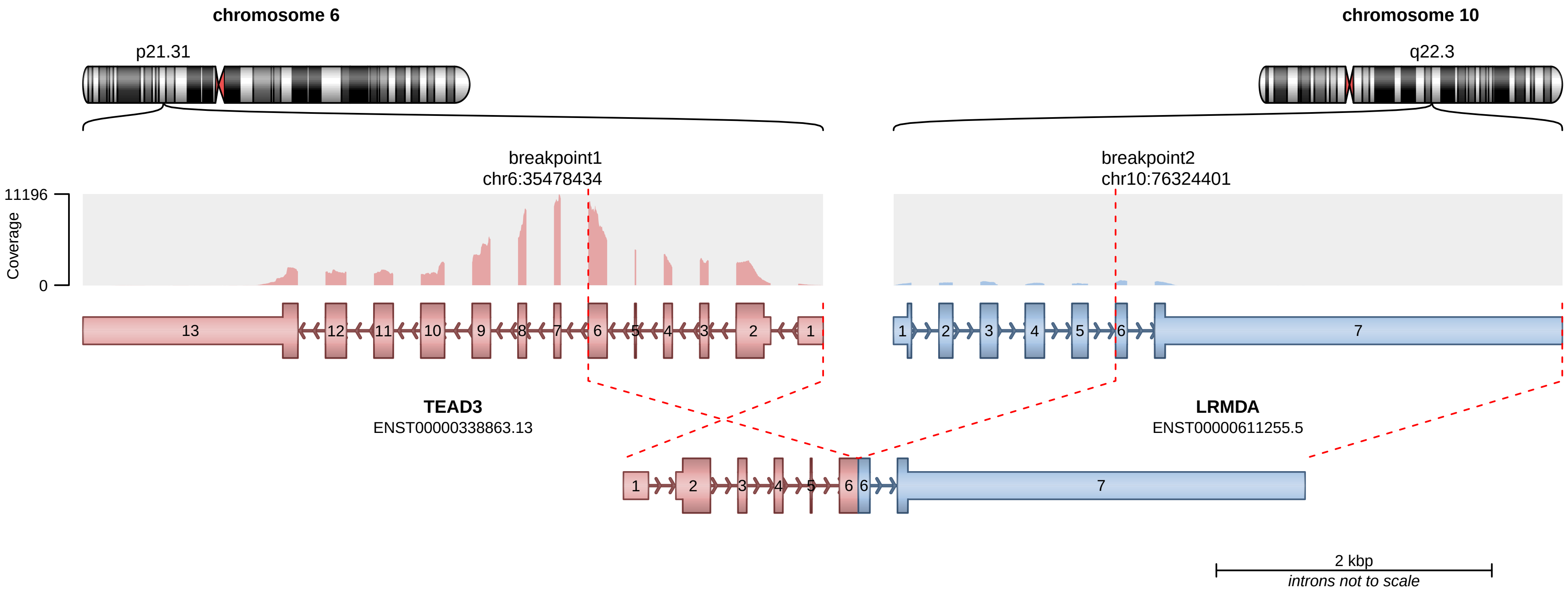


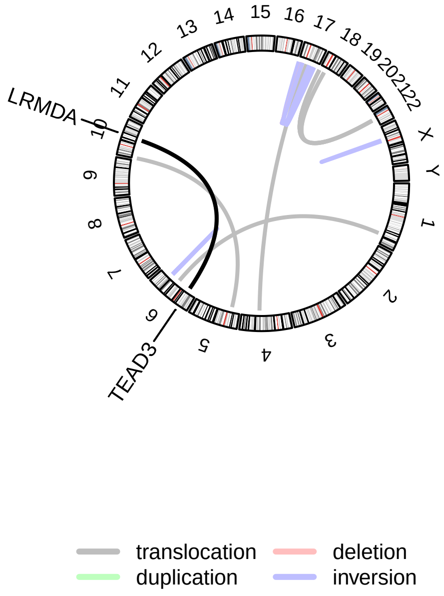


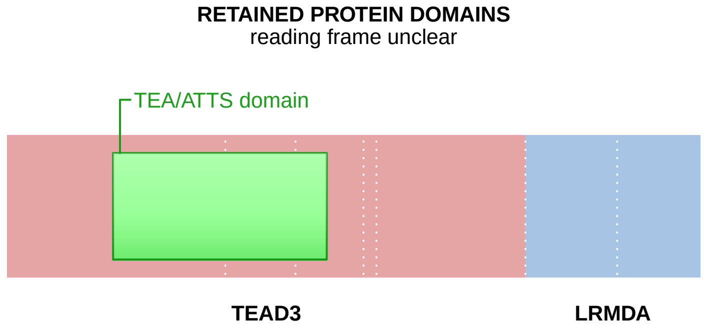


SUPPORTING READ COUNT

Split reads = 79
Discordant mates = 0

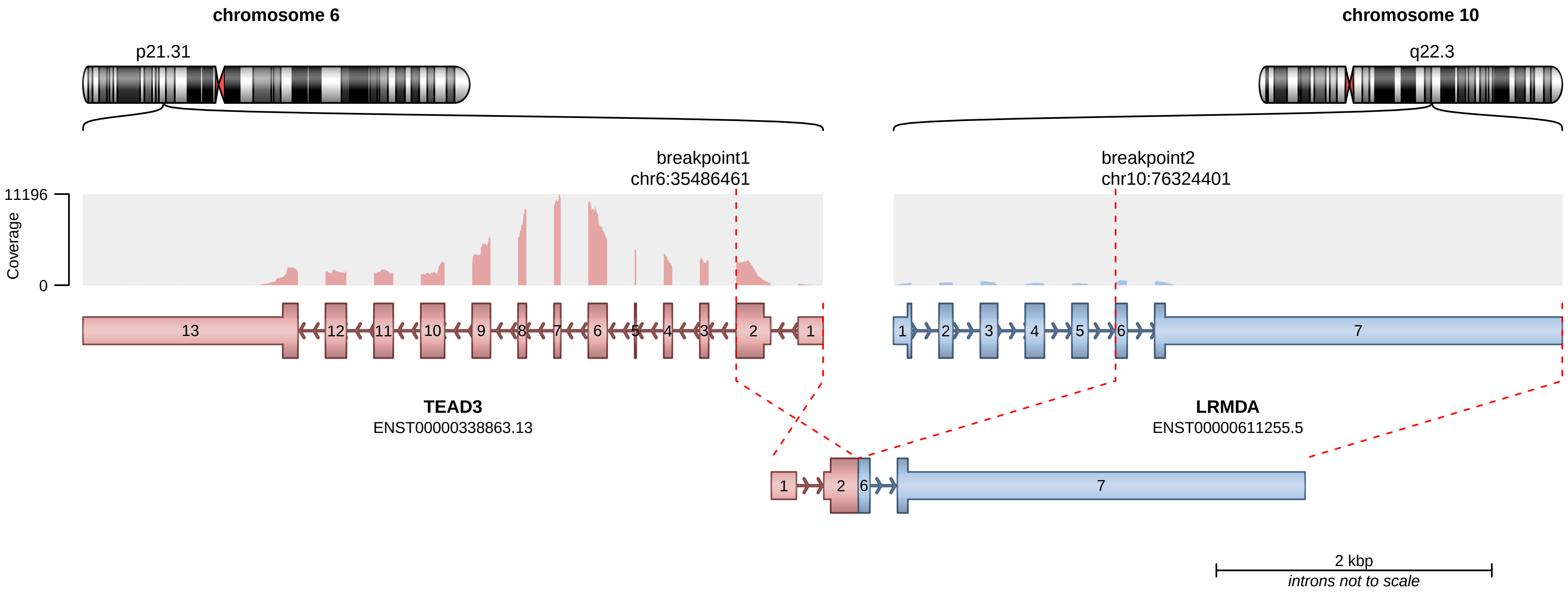


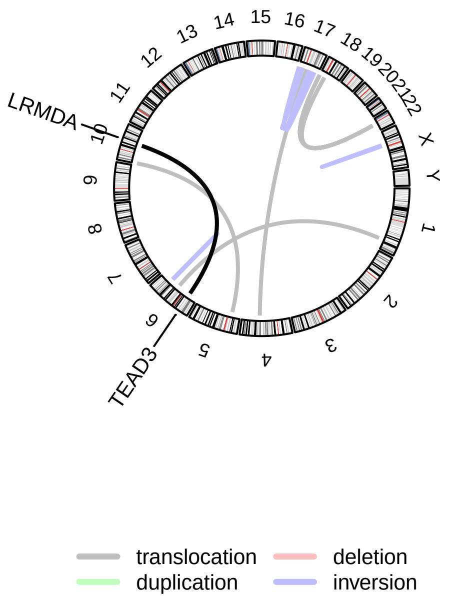


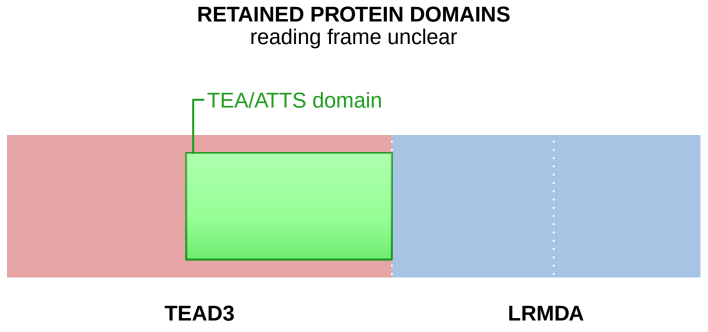


SUPPORTING READ COUNT

Split reads = 6
Discordant mates = 0

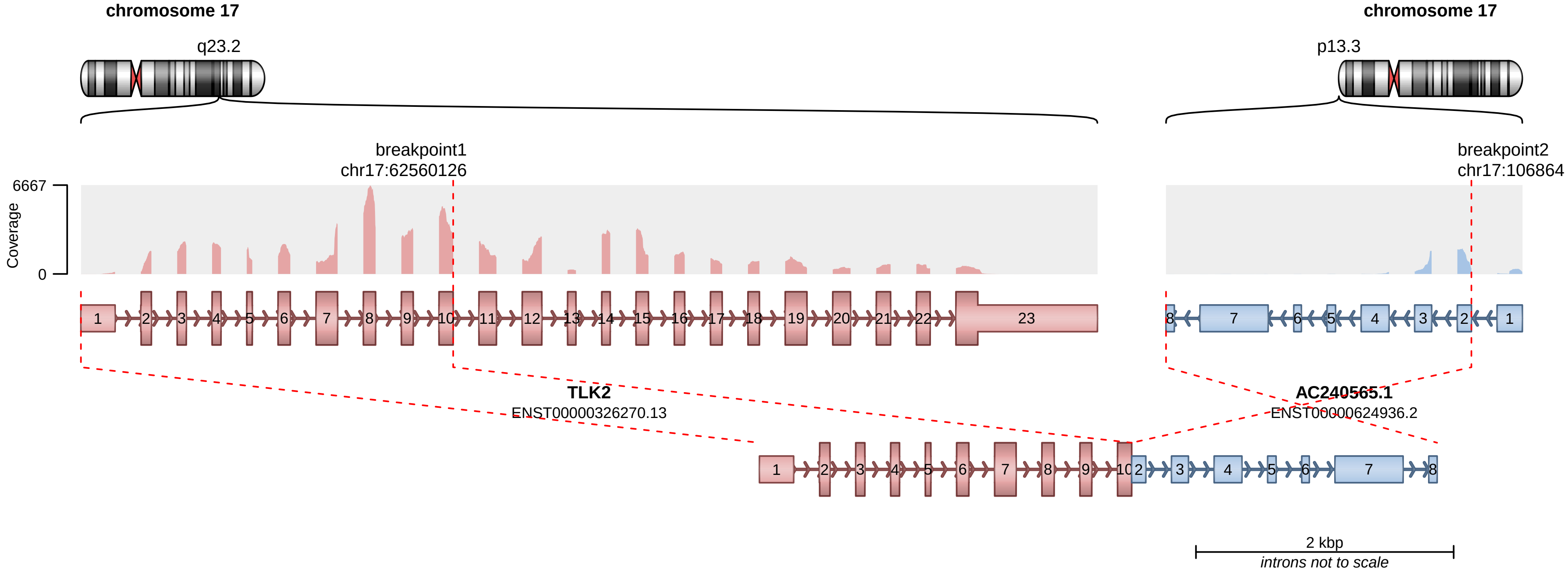


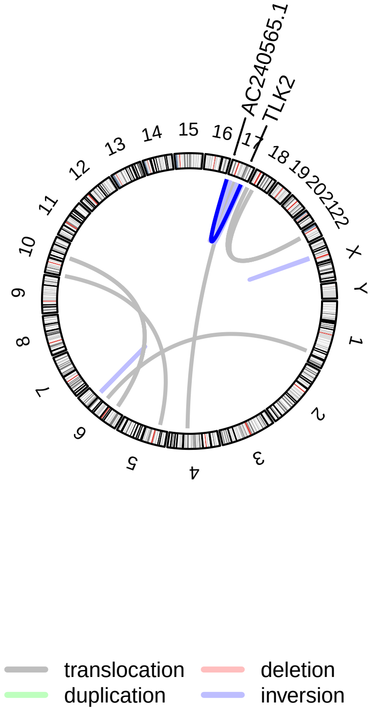




SUPPORTING READ COUNT

Split reads = 5
Discordant mates = 0

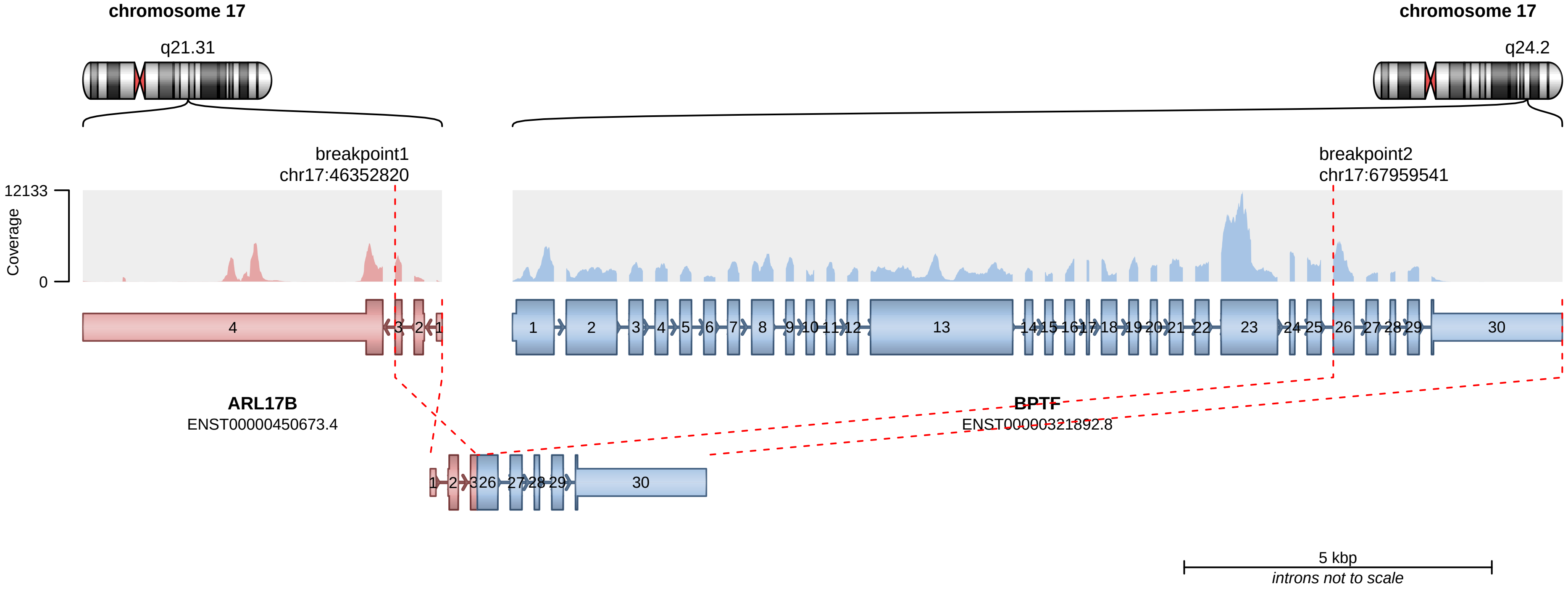


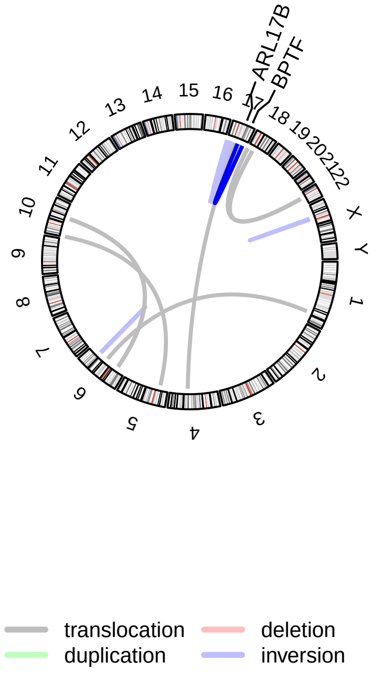


No protein domains retained in fusion.

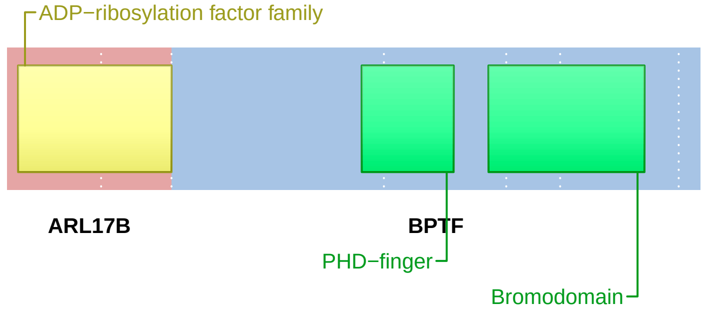
SUPPORTING READ COUNT

Split reads = 72
Discordant mates = 0



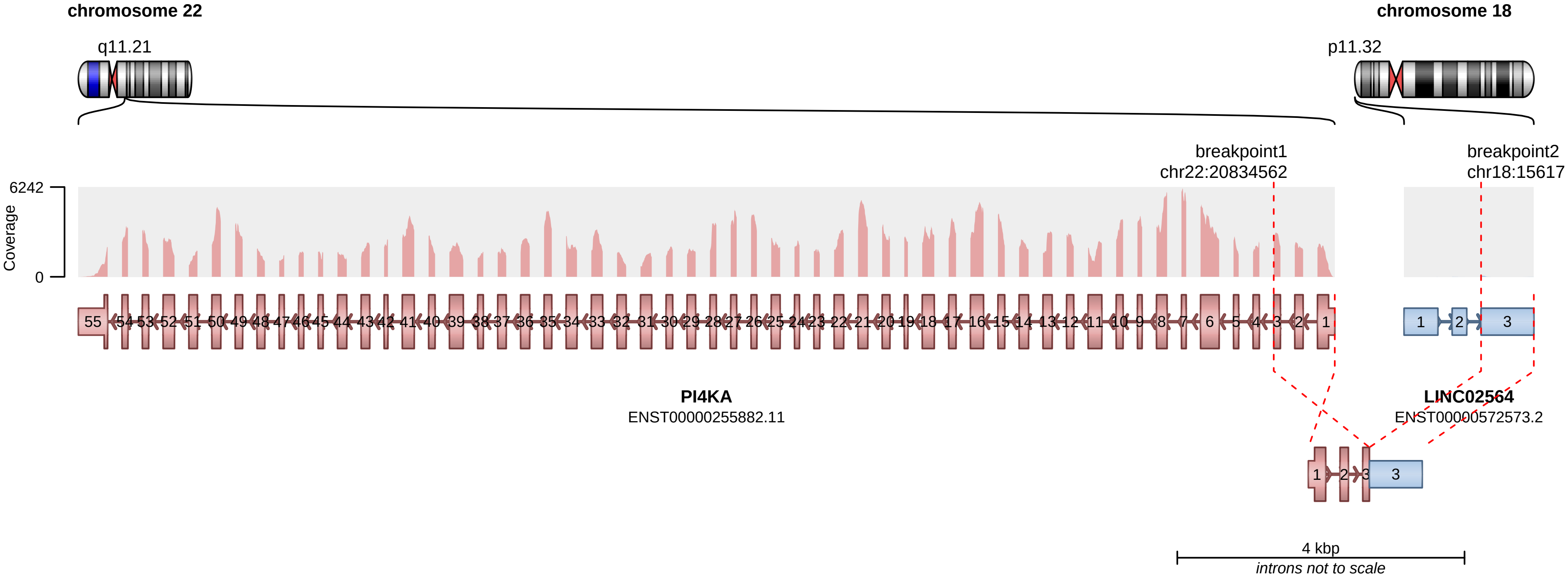


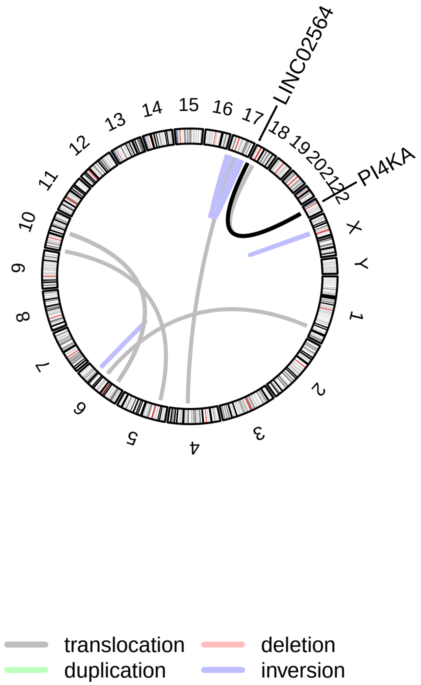
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 62
Discordant mates = 0

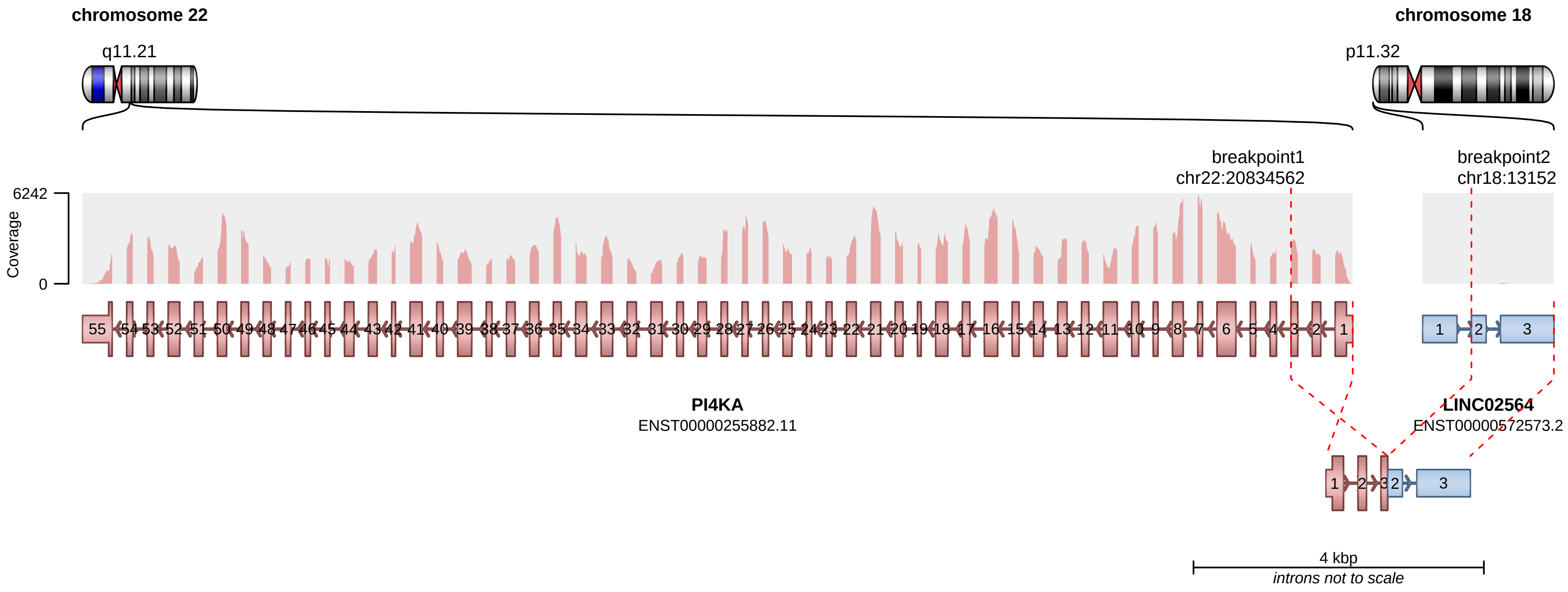


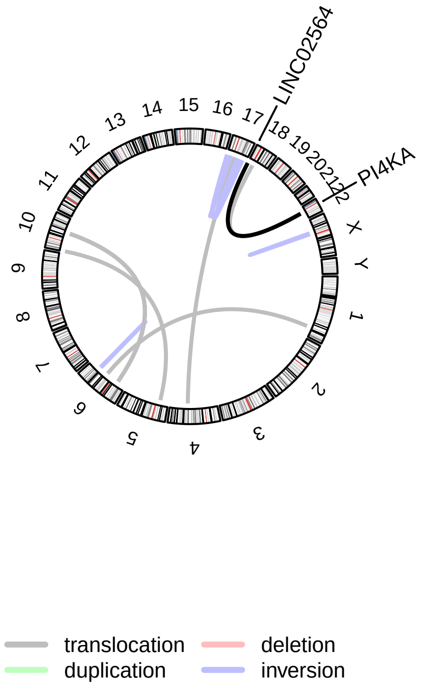


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 53
Discordant mates = 0

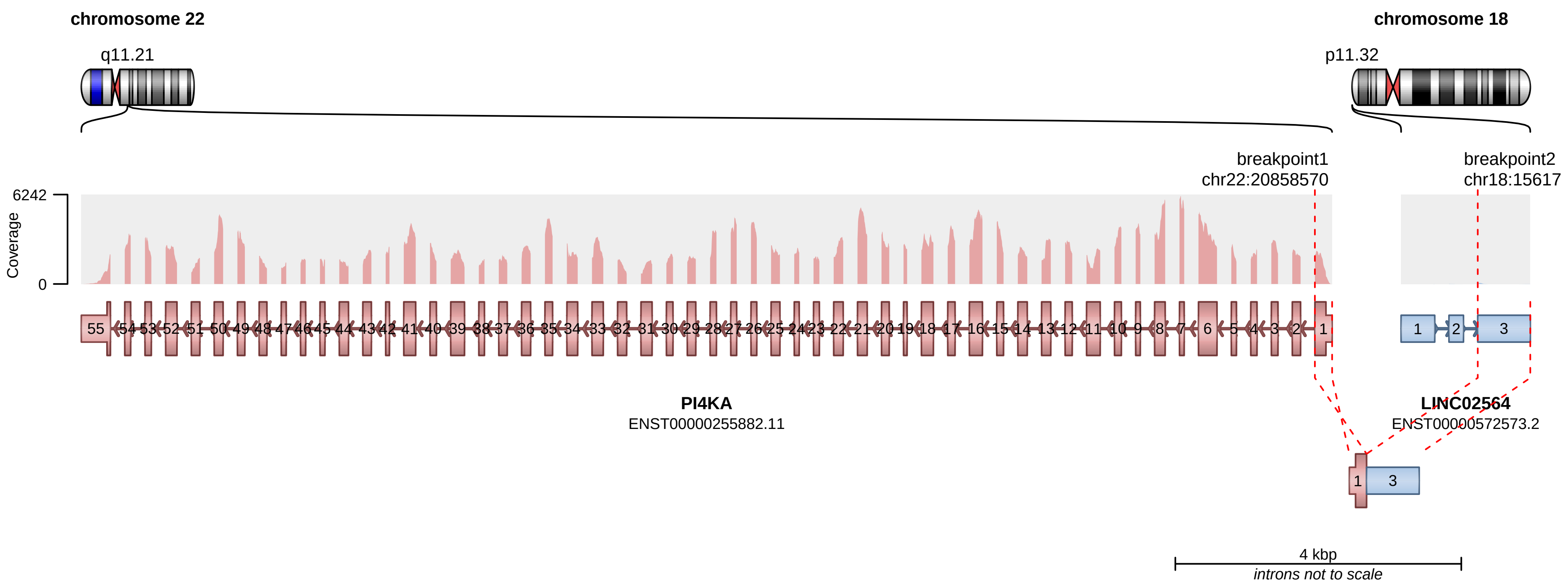


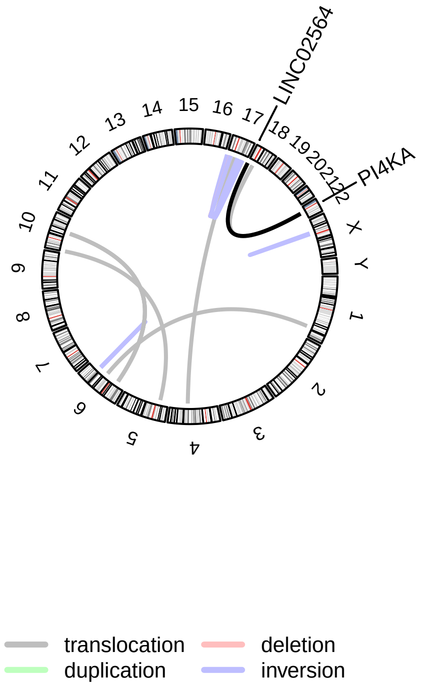


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 0

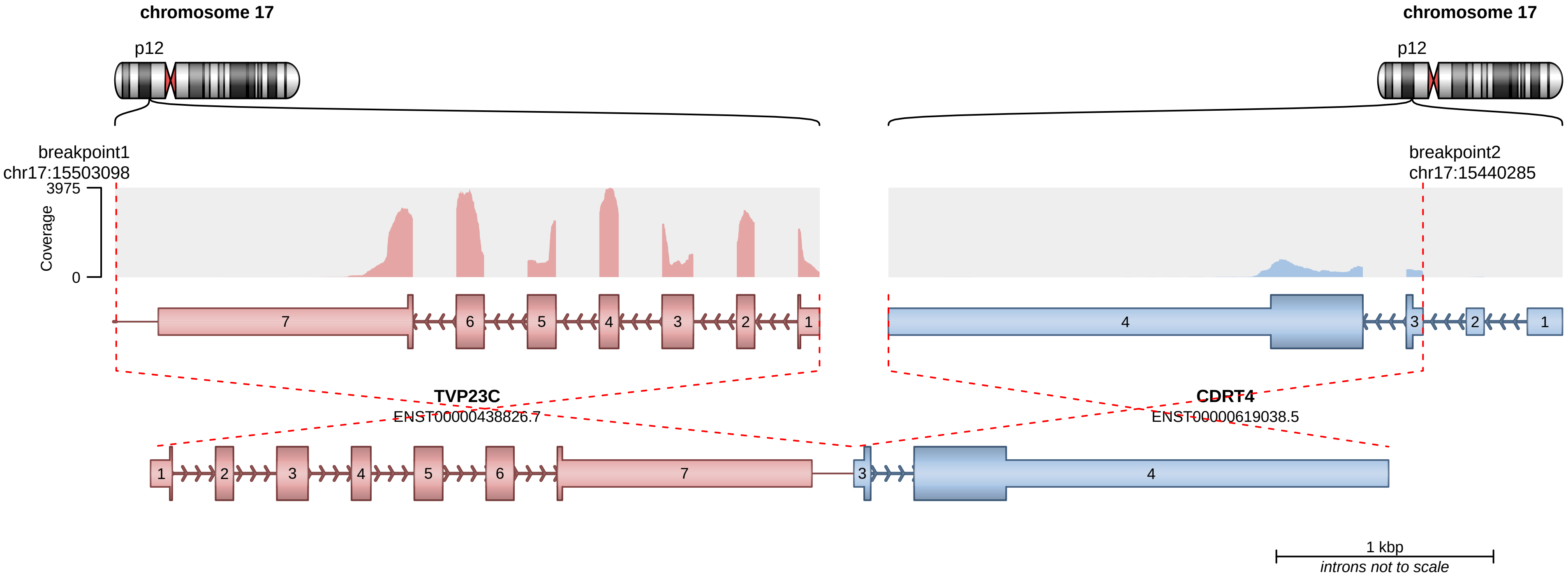


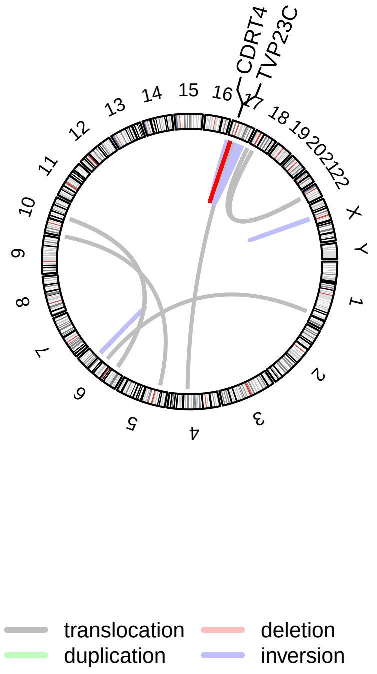


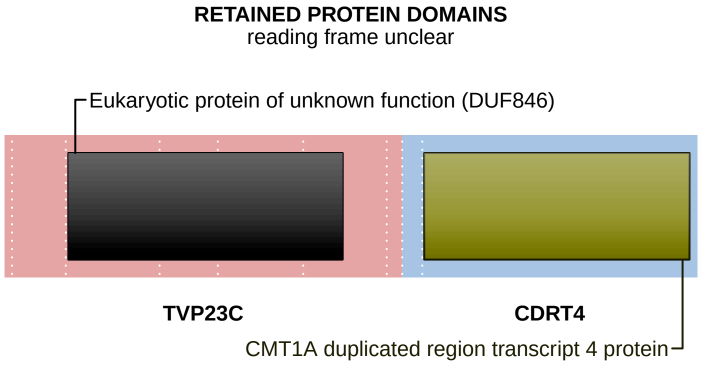
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

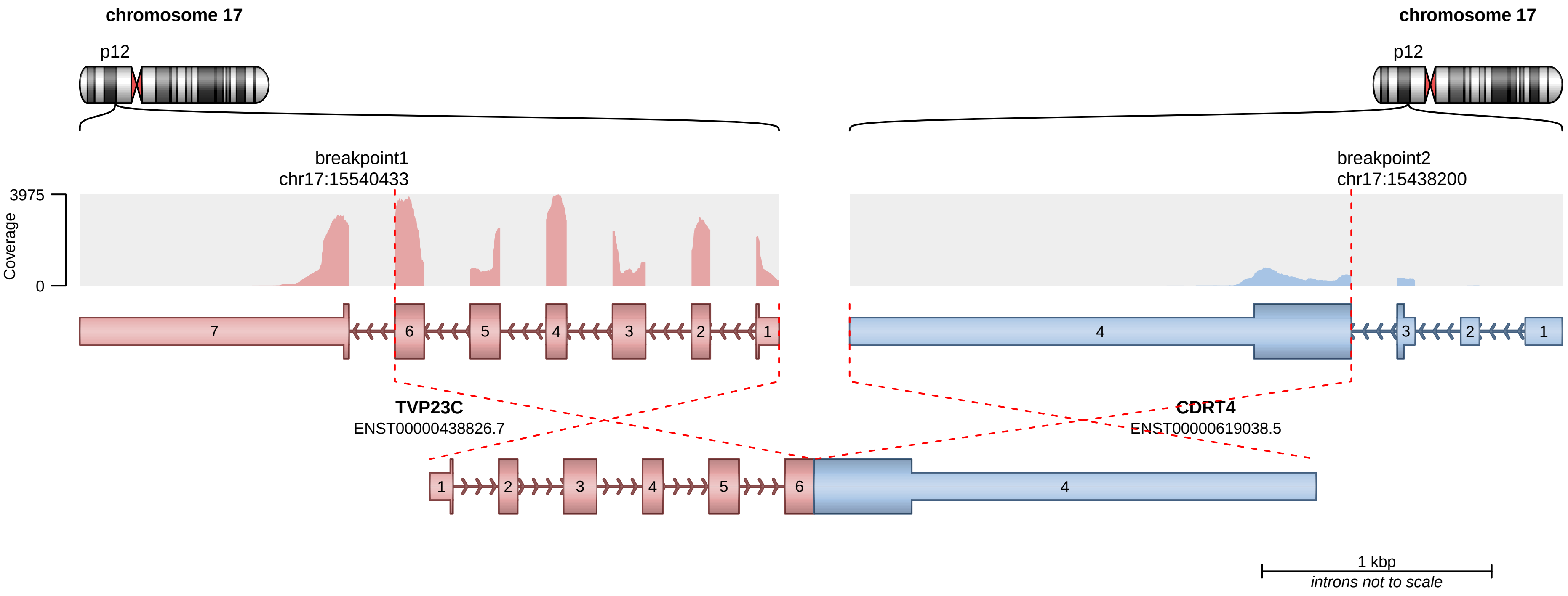


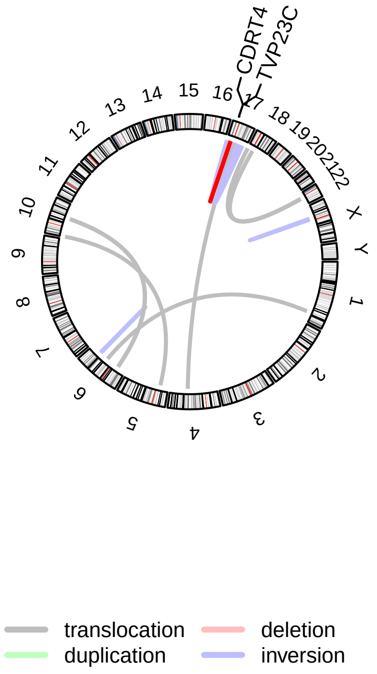




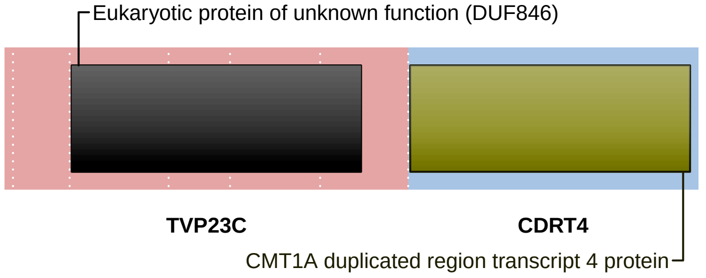
SUPPORTING READ COUNT

Split reads = 47
Discordant mates = 0



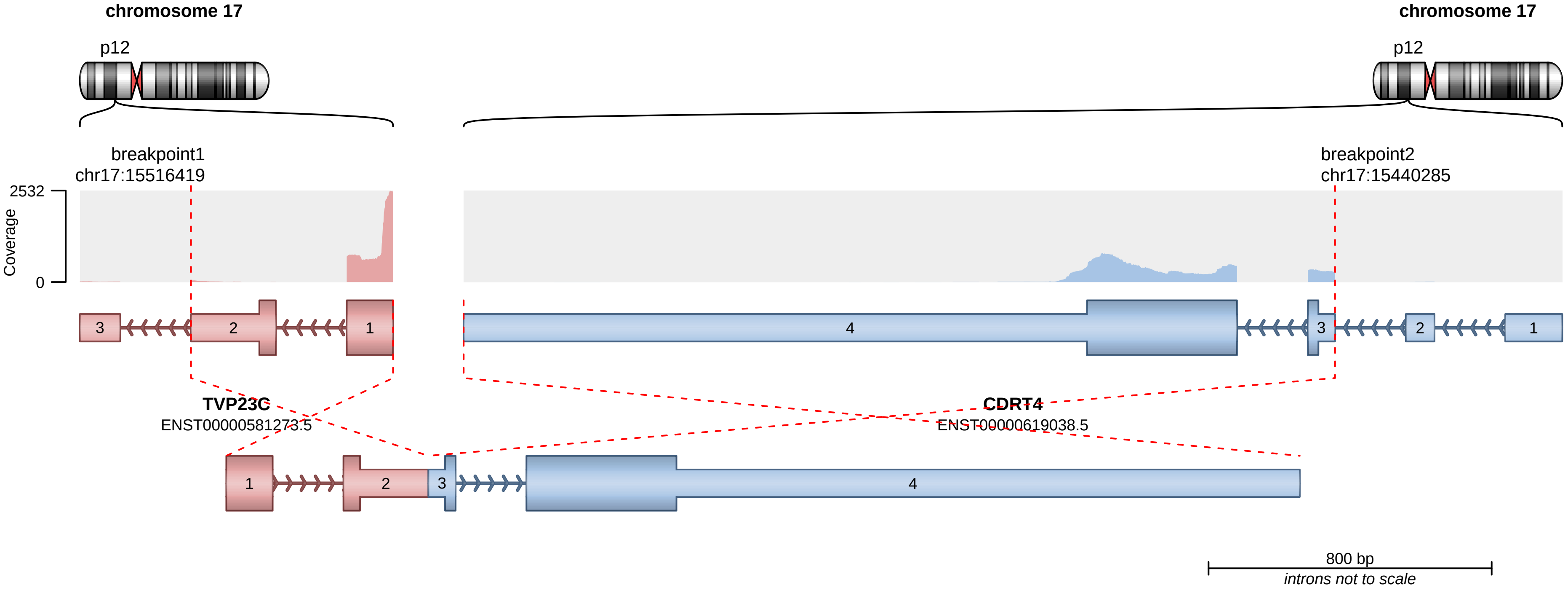


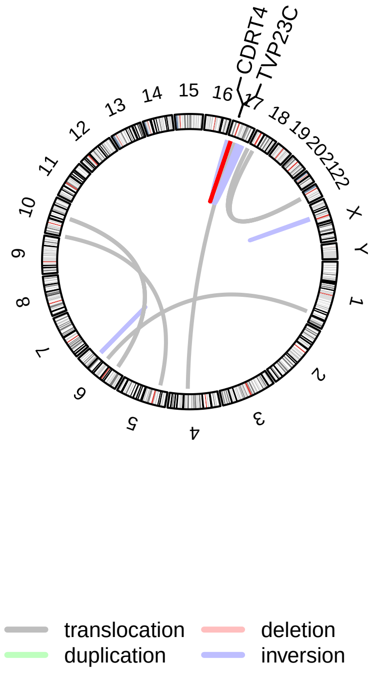
RETAINED PROTEIN DOMAINS
reading frame unclear

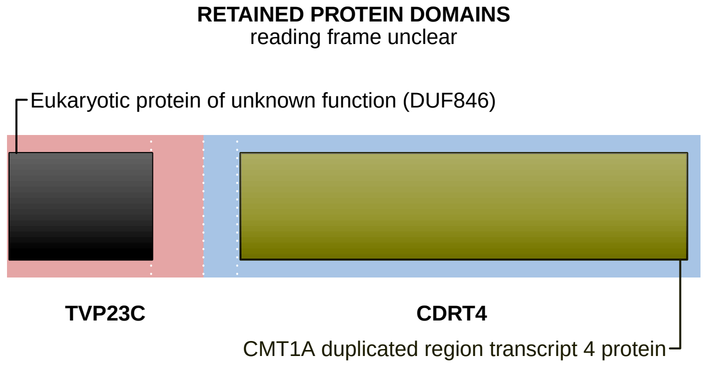


SUPPORTING READ COUNT

Split reads = 12
Discordant mates = 0

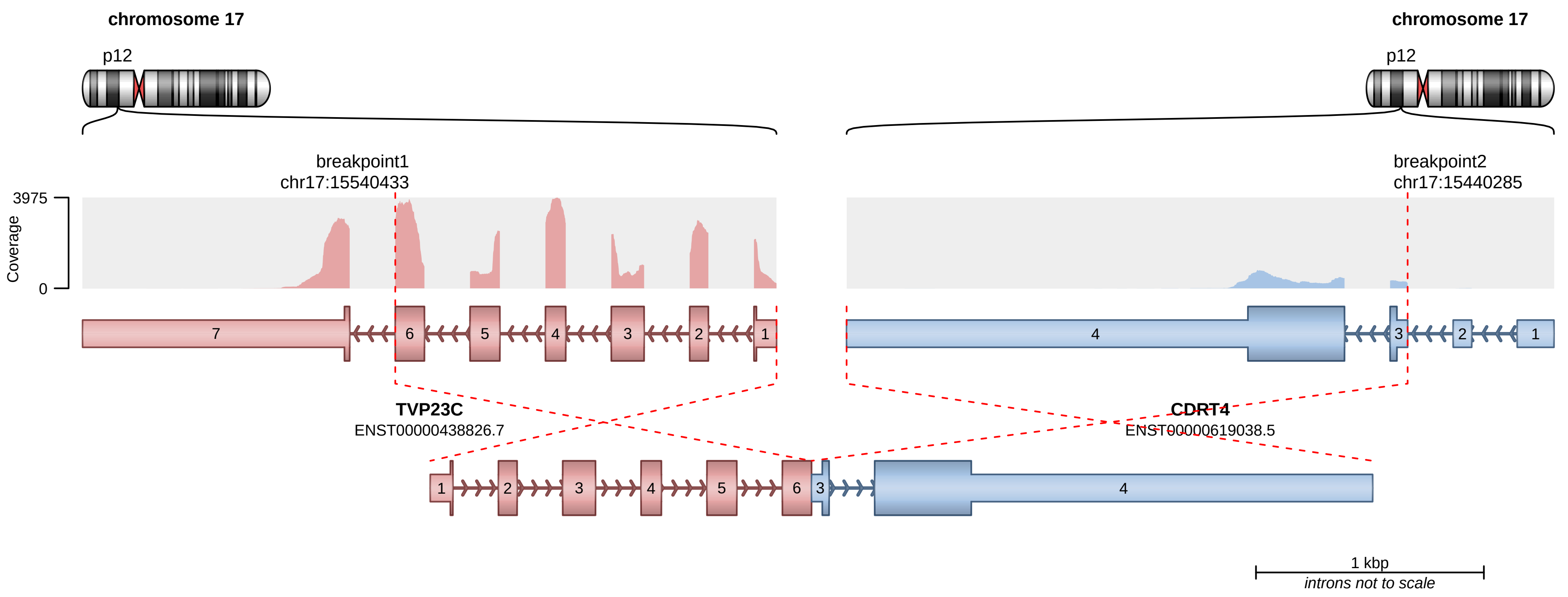


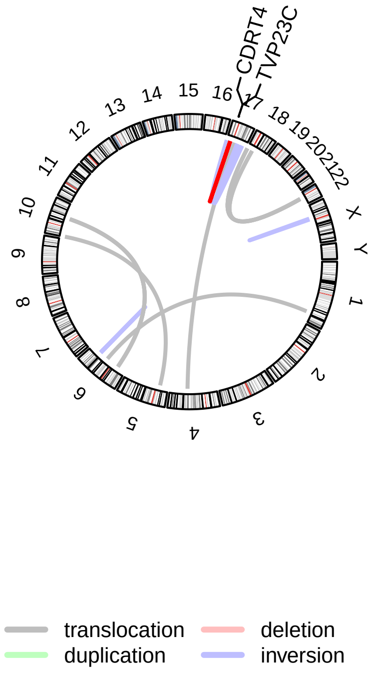




SUPPORTING READ COUNT

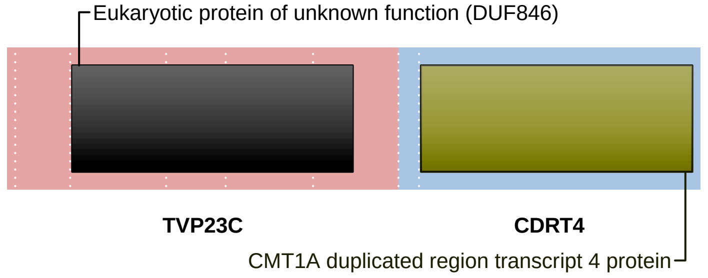
Split reads = 9
Discordant mates = 0





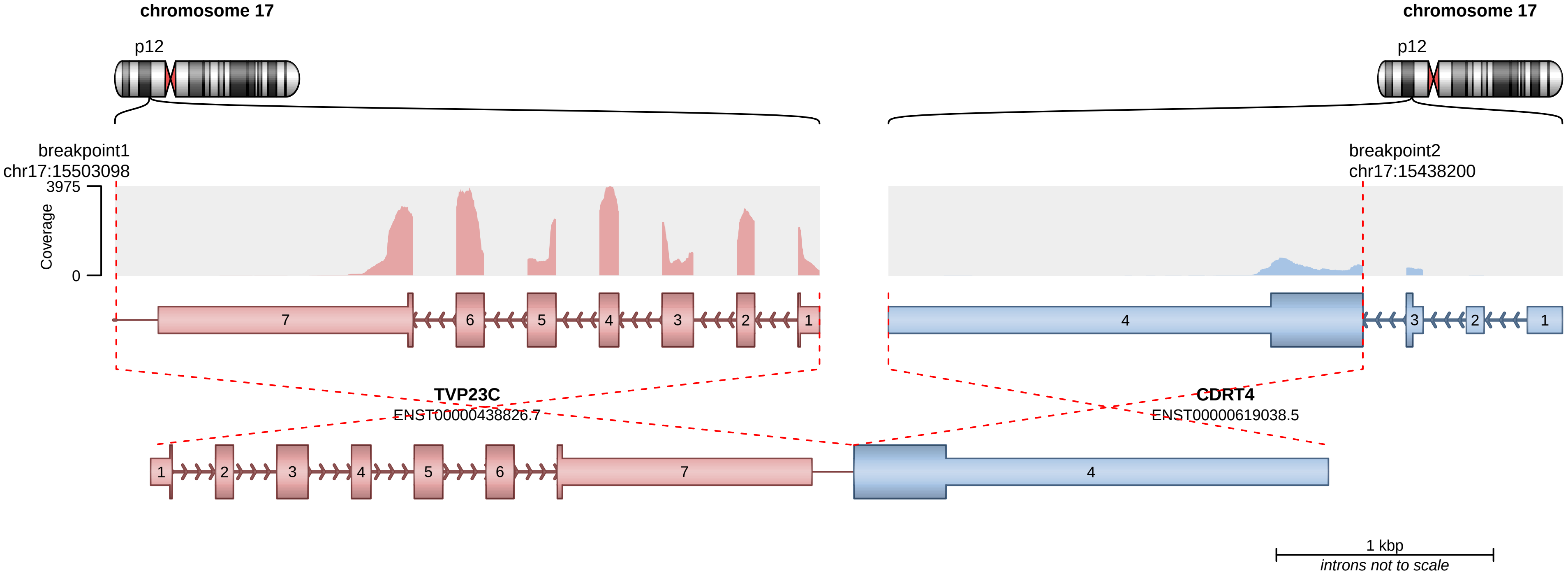
RETAINED PROTEIN DOMAINS

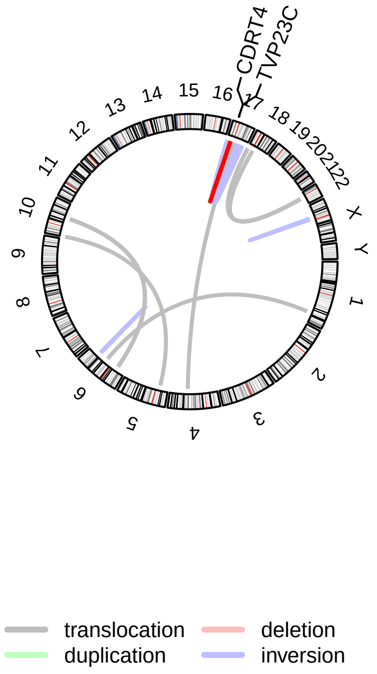
reading frame unclear

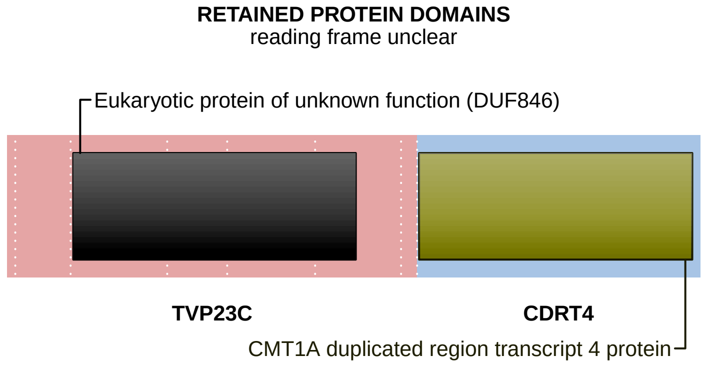


SUPPORTING READ COUNT

Split reads = 8
Discordant mates = 0



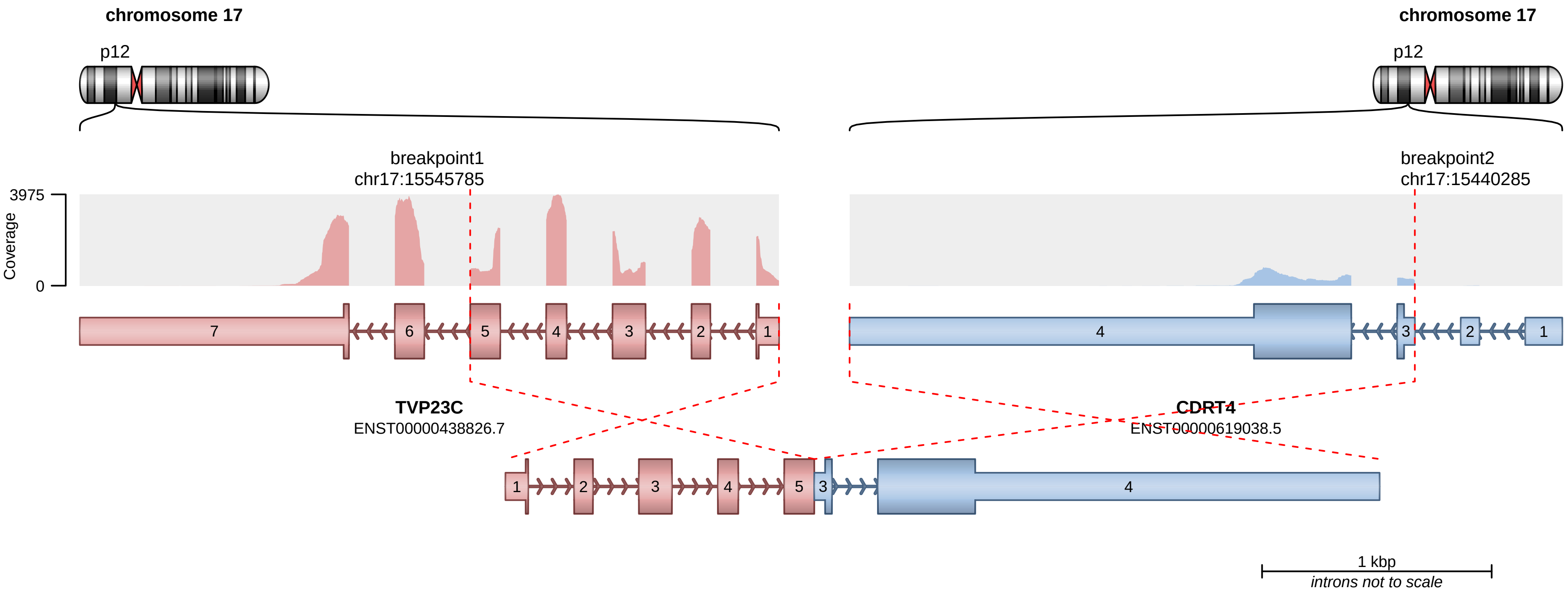


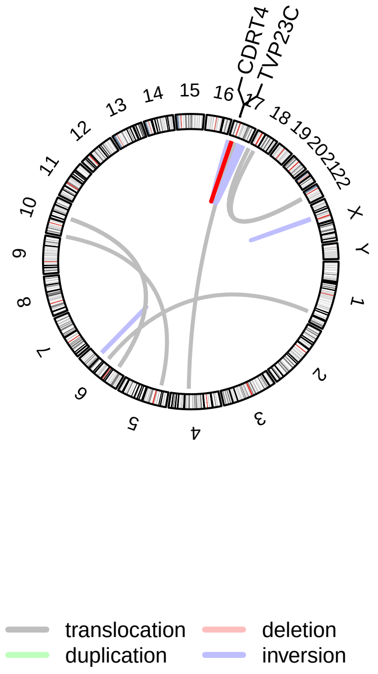

RETAINED PROTEIN DOMAINS

reading frame unclear

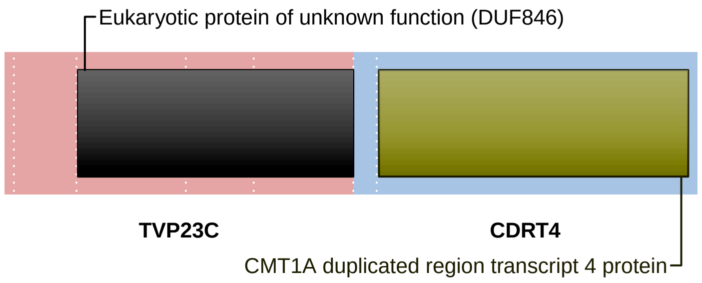
SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0



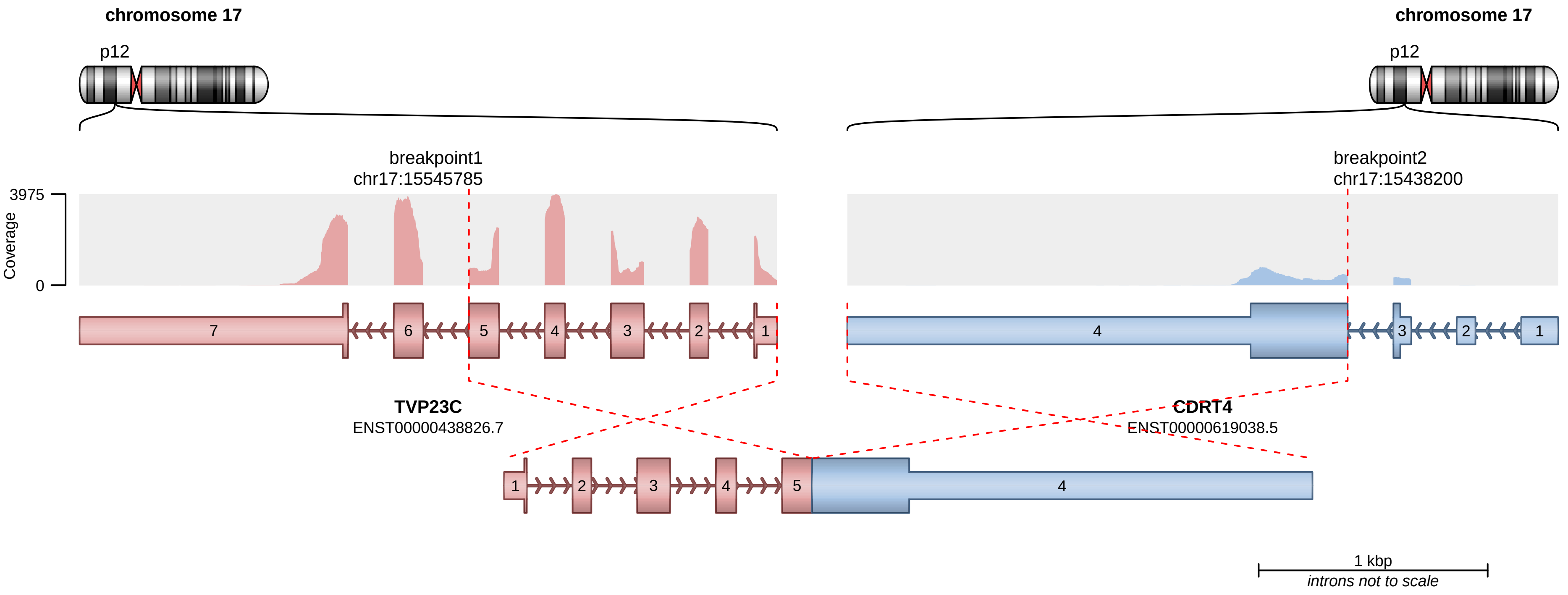


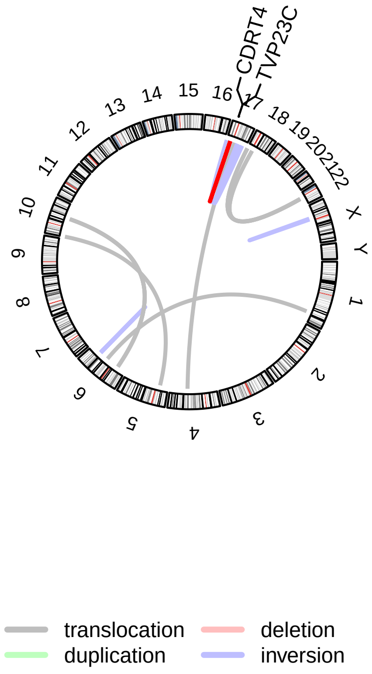
RETAINED PROTEIN DOMAINS
reading frame unclear

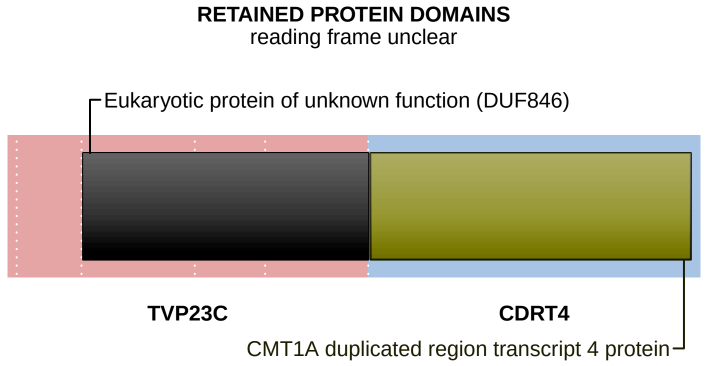


SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

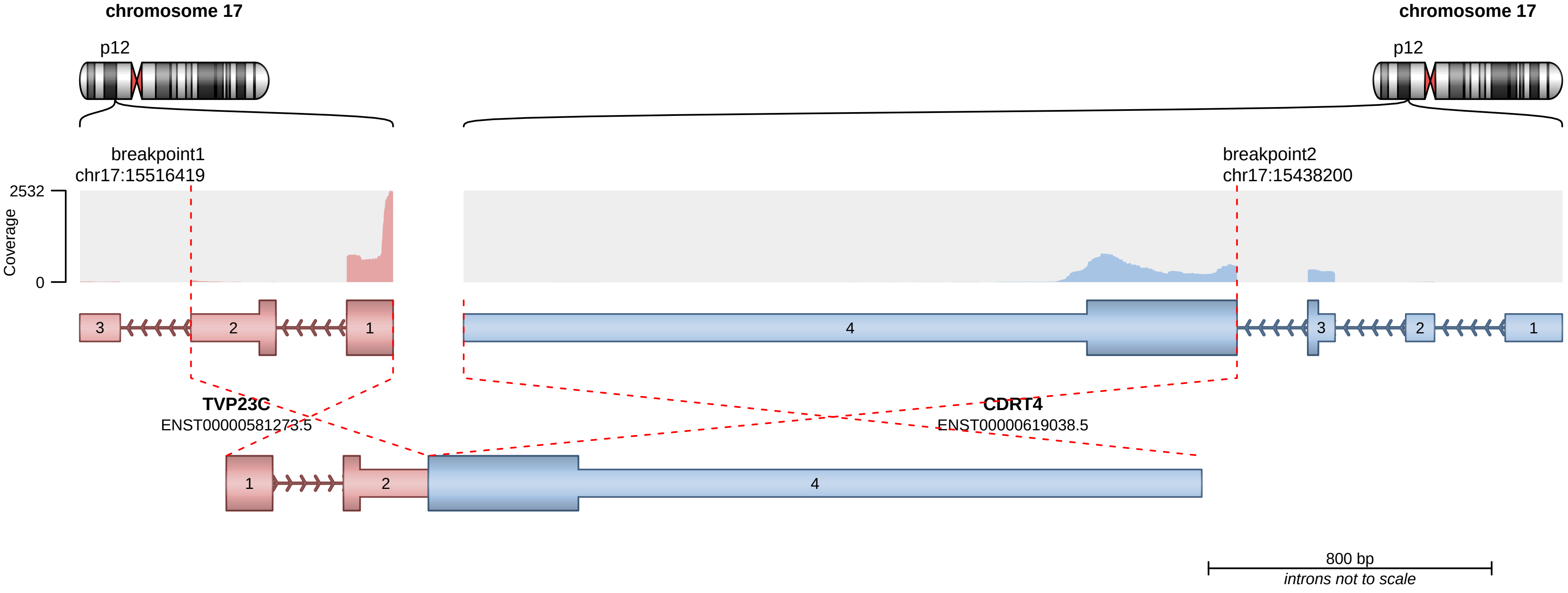


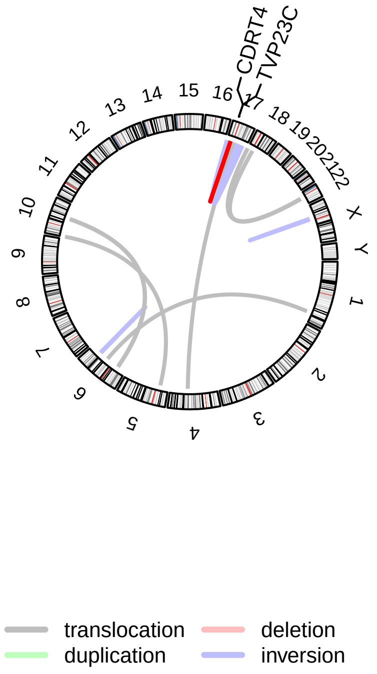


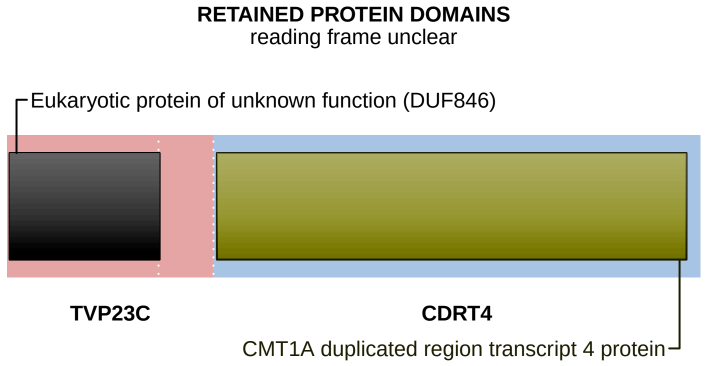


SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

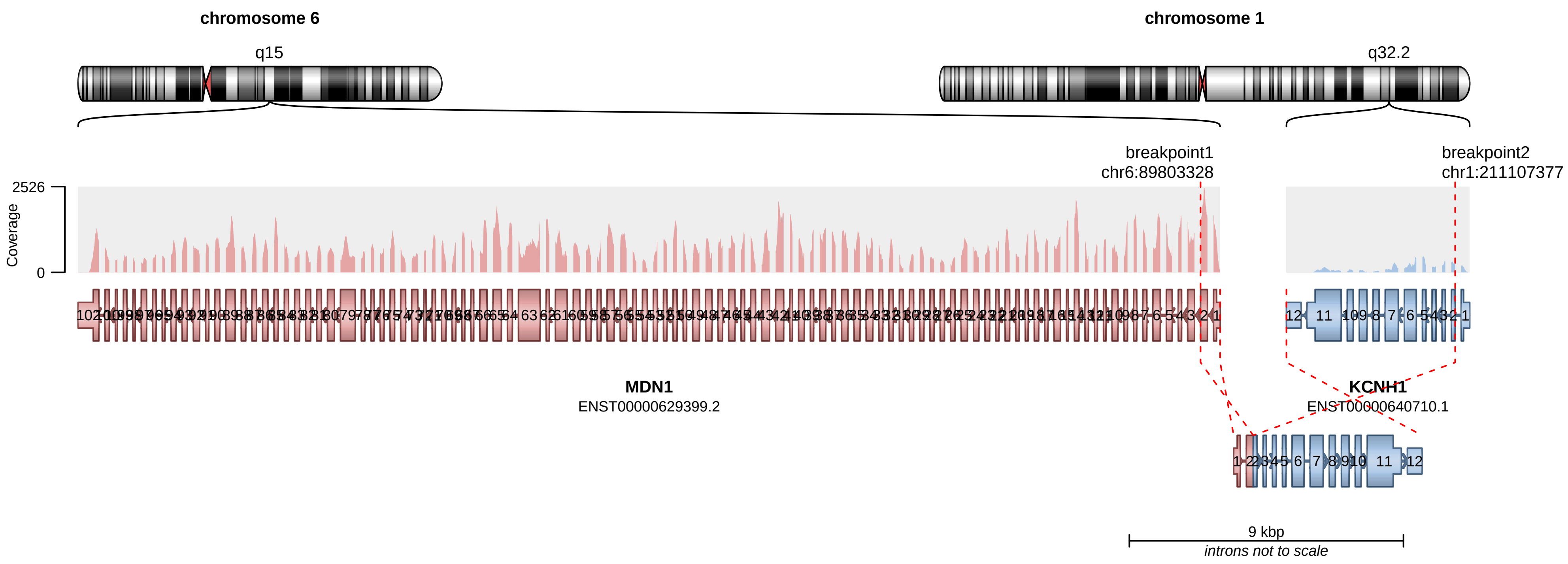


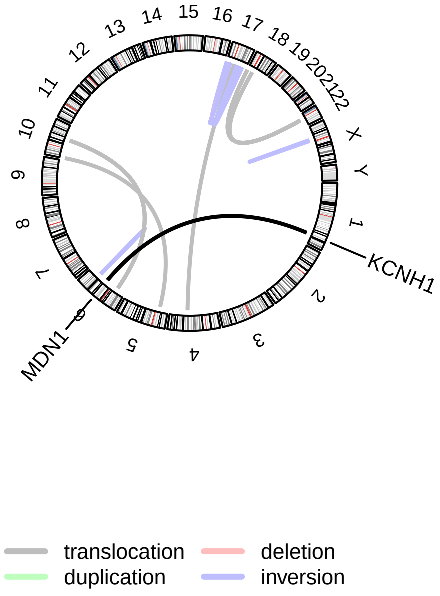




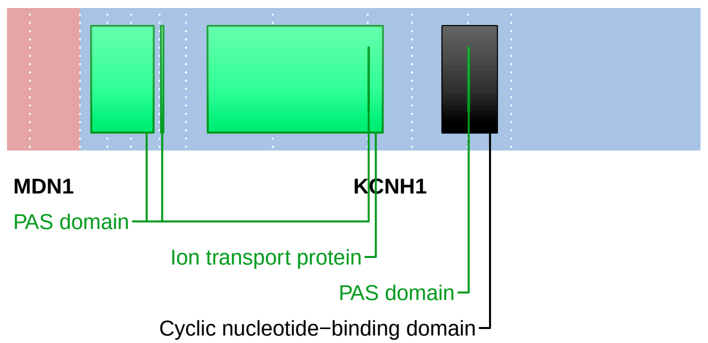
SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0



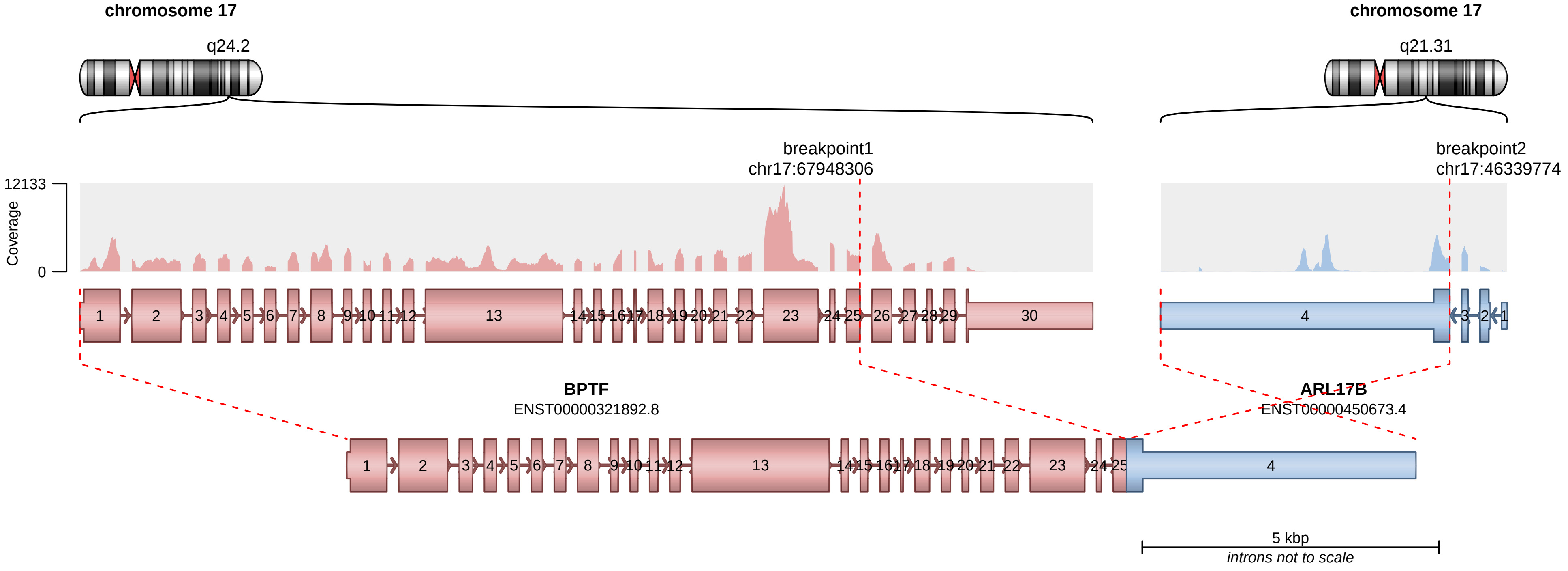


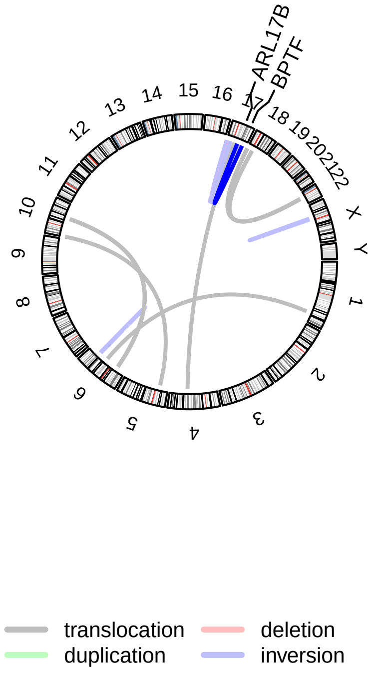
RETAINED PROTEIN DOMAINS
reading frame unclear



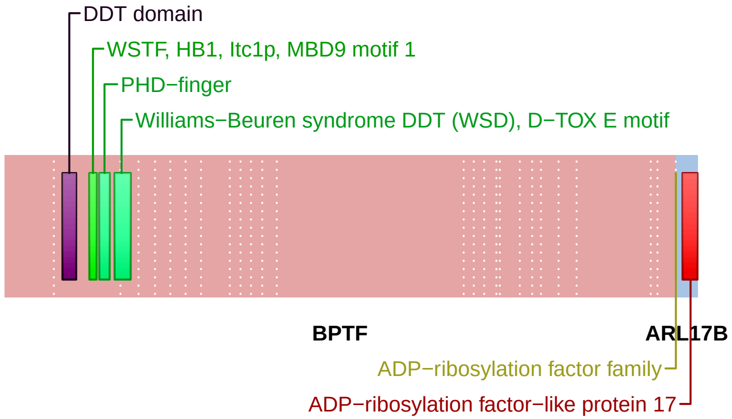
SUPPORTING READ COUNT

Split reads = 30
Discordant mates = 0



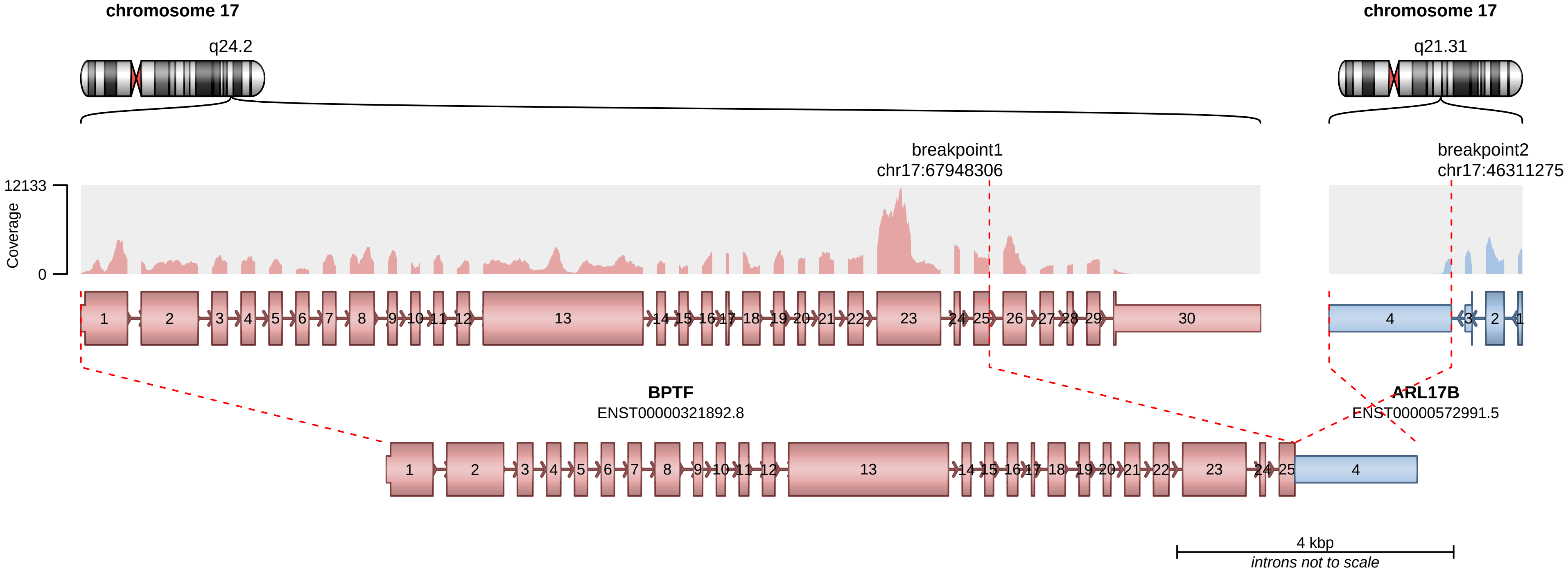


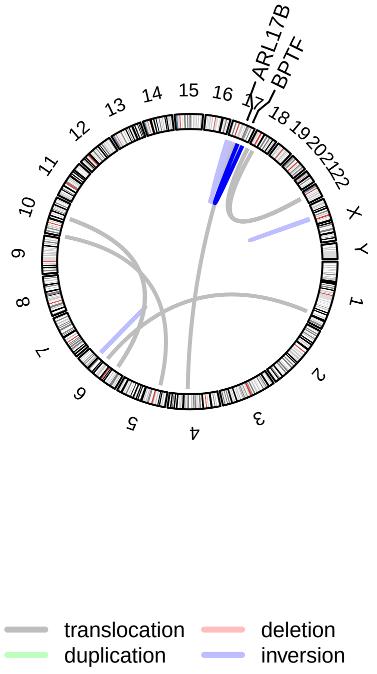
RETAINED PROTEIN DOMAINS
reading frame unclear

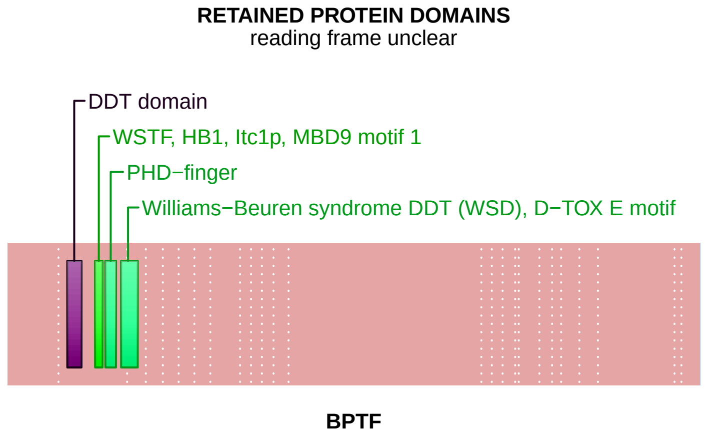


SUPPORTING READ COUNT

Split reads = 25
Discordant mates = 0

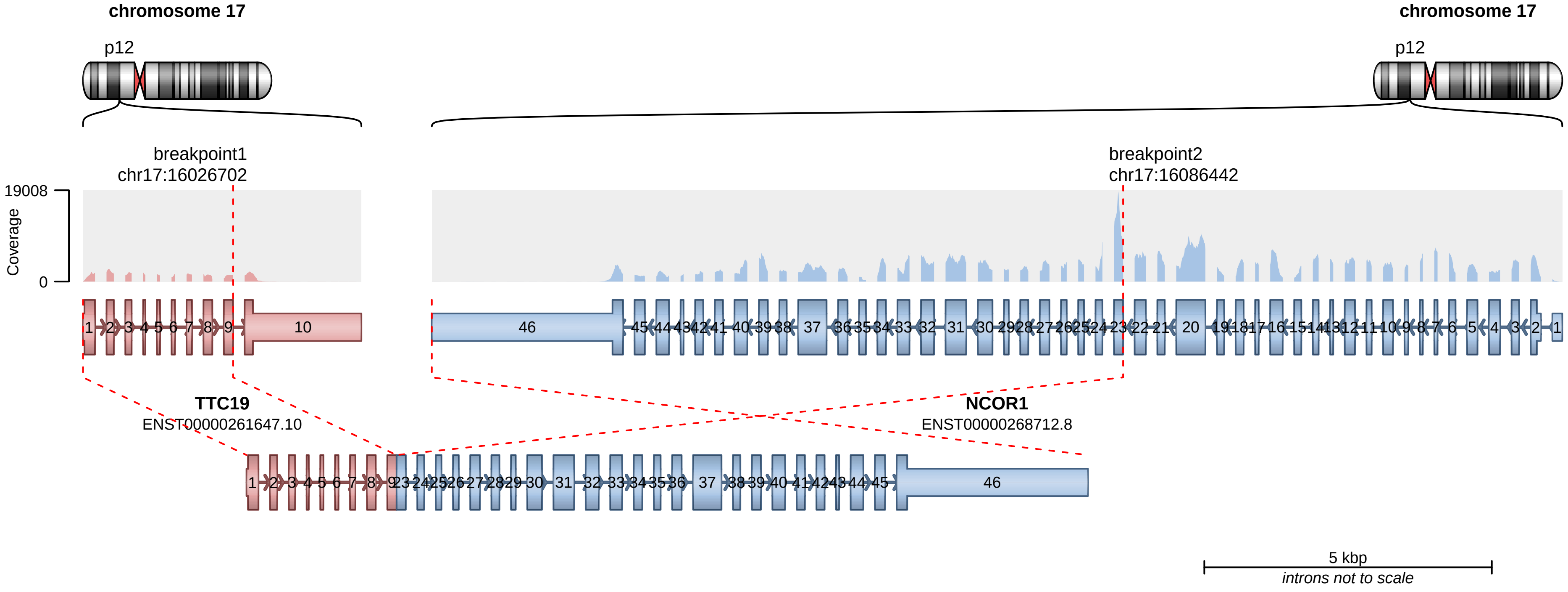


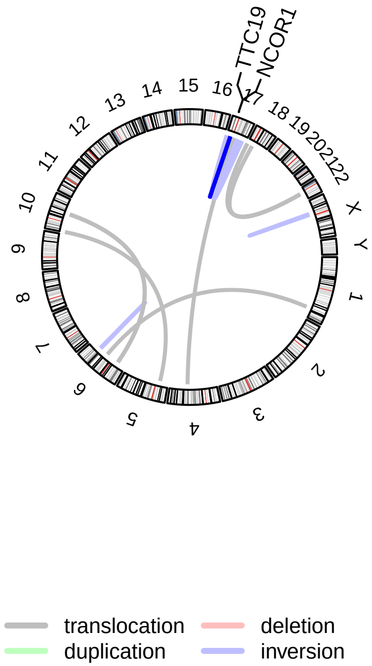


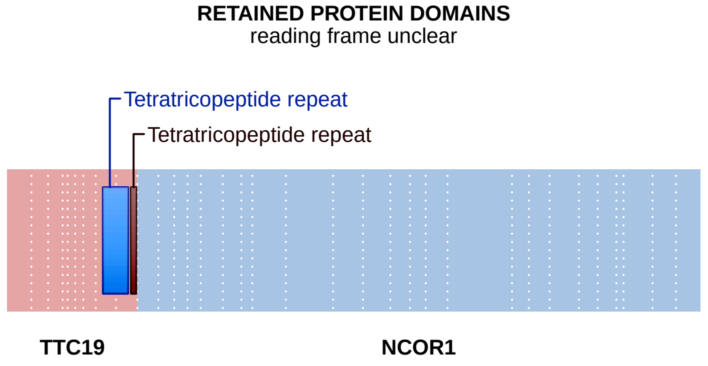


SUPPORTING READ COUNT

Split reads = 5
Discordant mates = 0

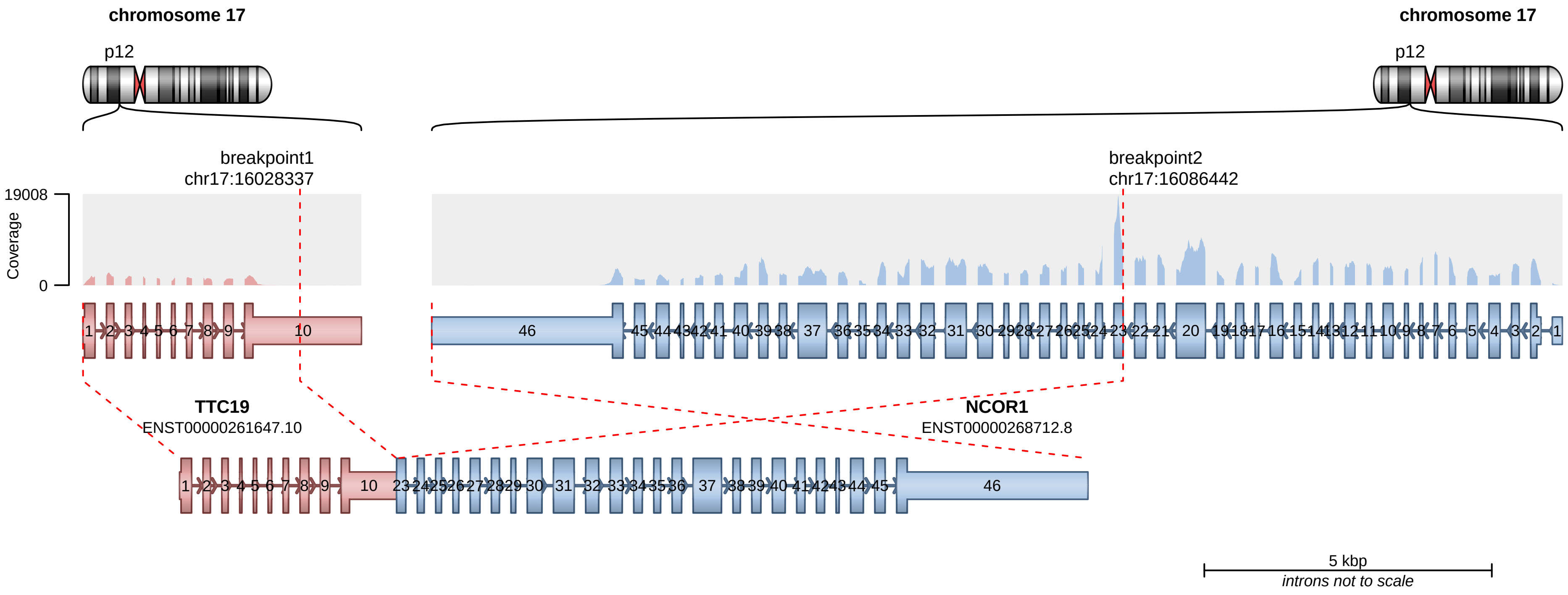


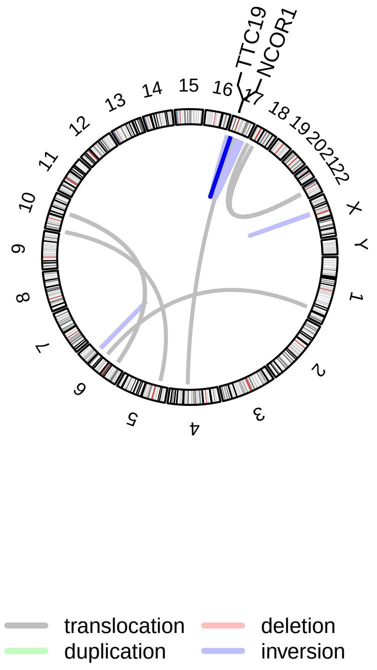




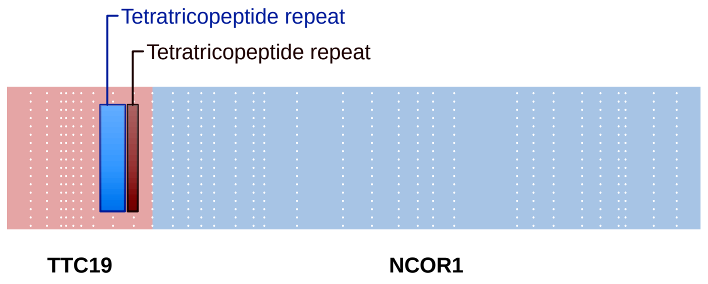
SUPPORTING READ COUNT

Split reads = 19
Discordant mates = 0



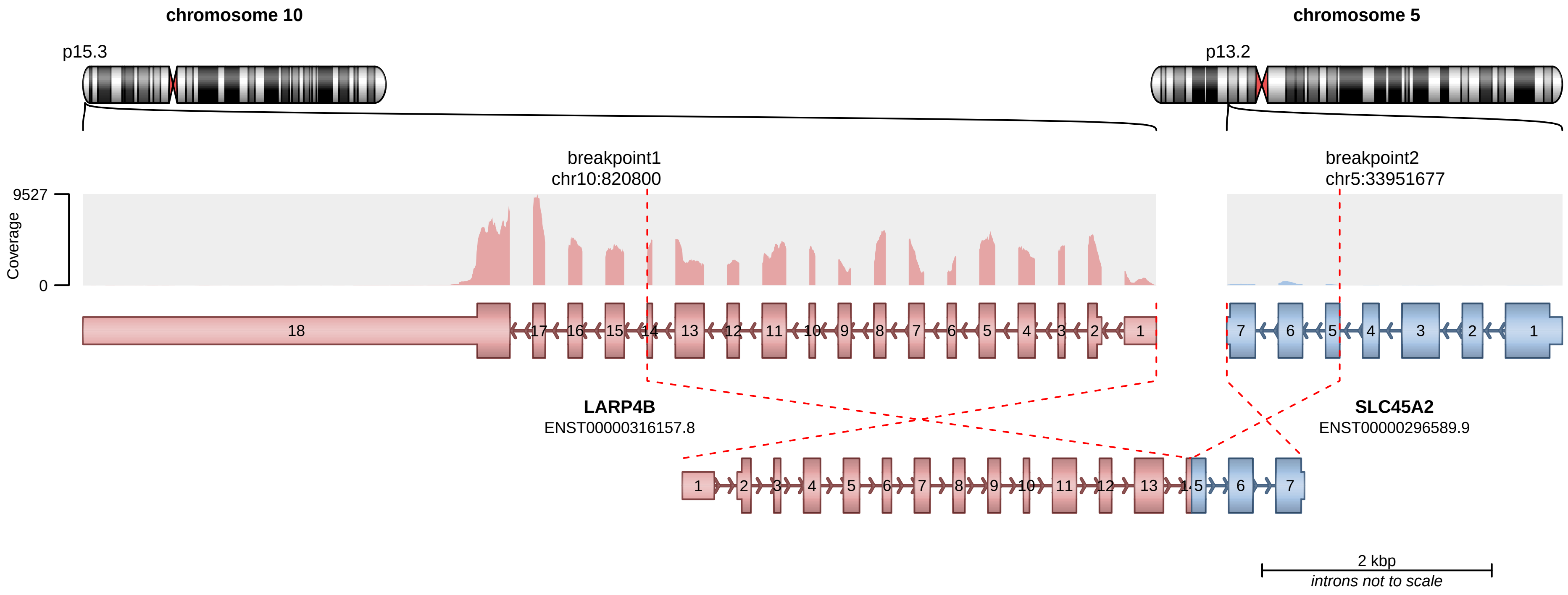


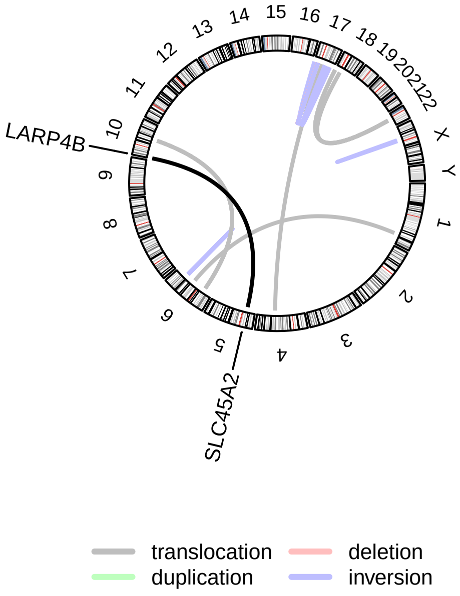
RETAINED PROTEIN DOMAINS
reading frame unclear



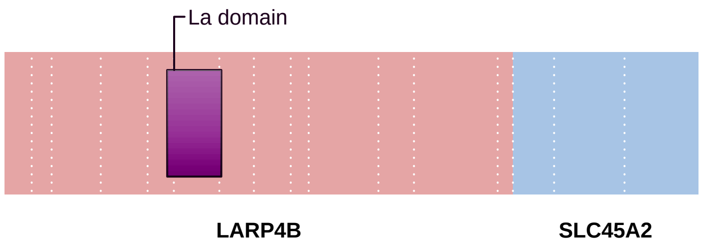
SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0



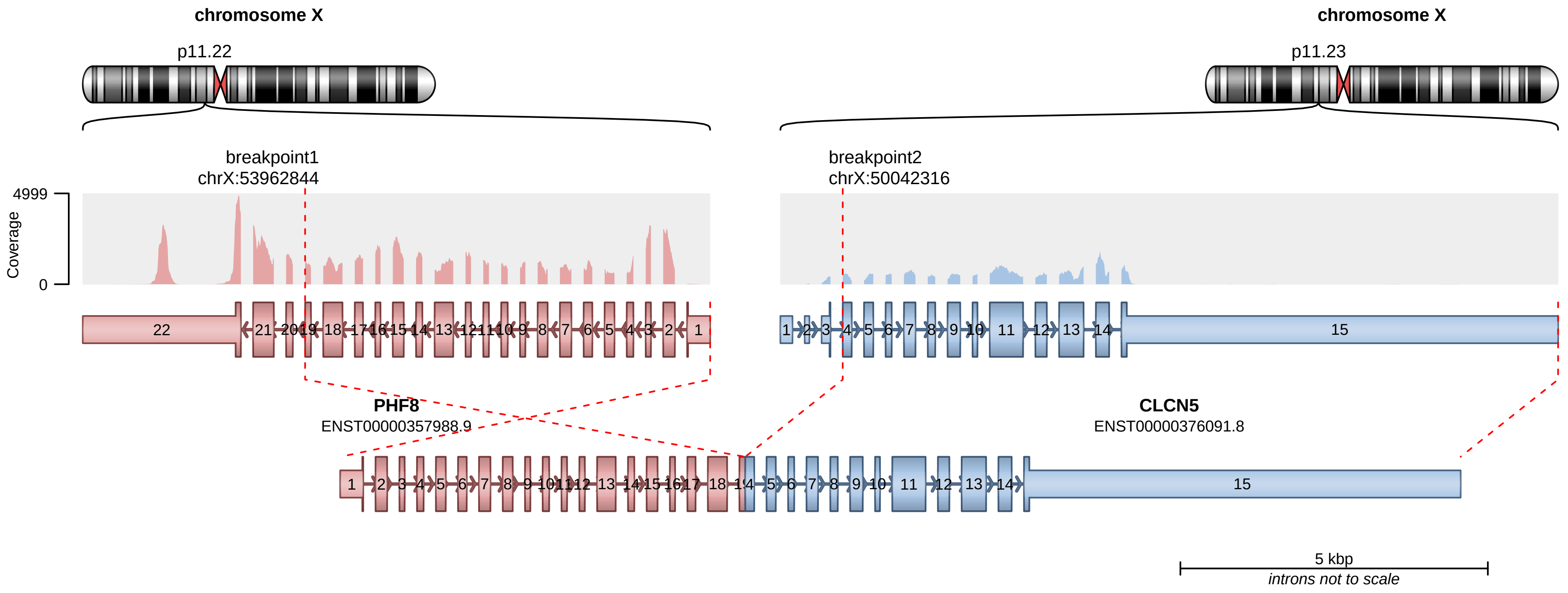


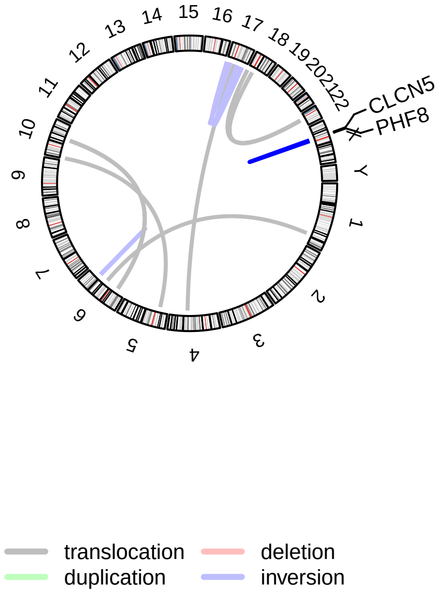
RETAINED PROTEIN DOMAINS
reading frame unclear

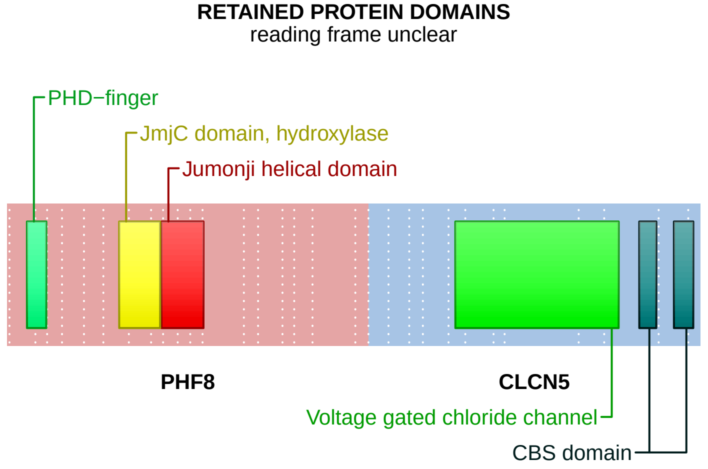


SUPPORTING READ COUNT

Split reads = 13
Discordant mates = 0







SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 0