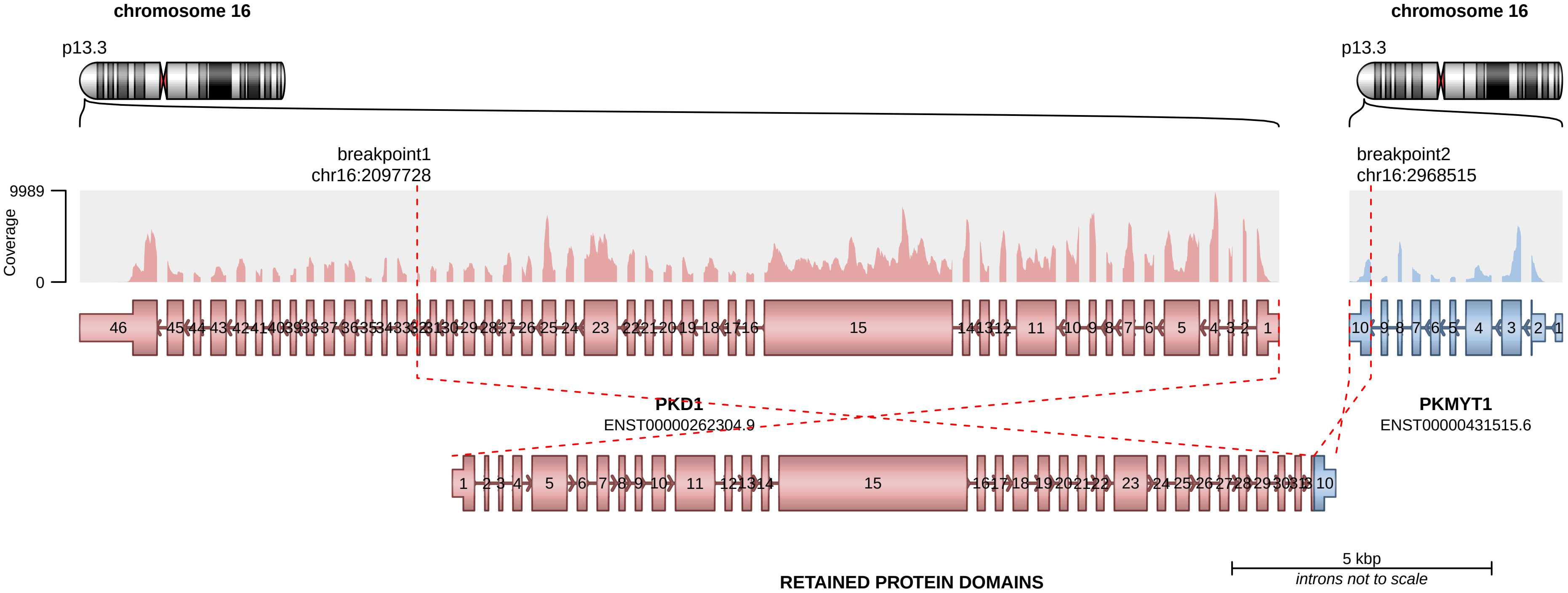
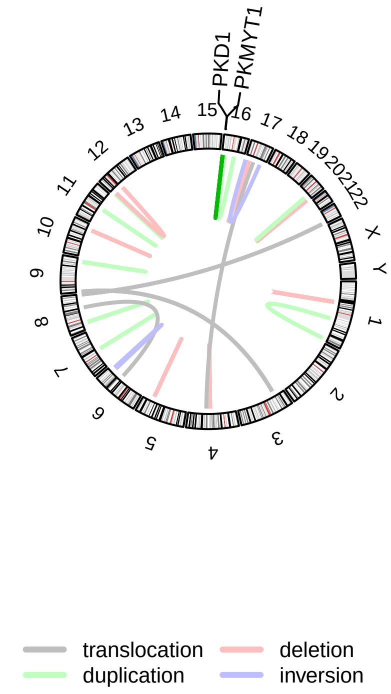
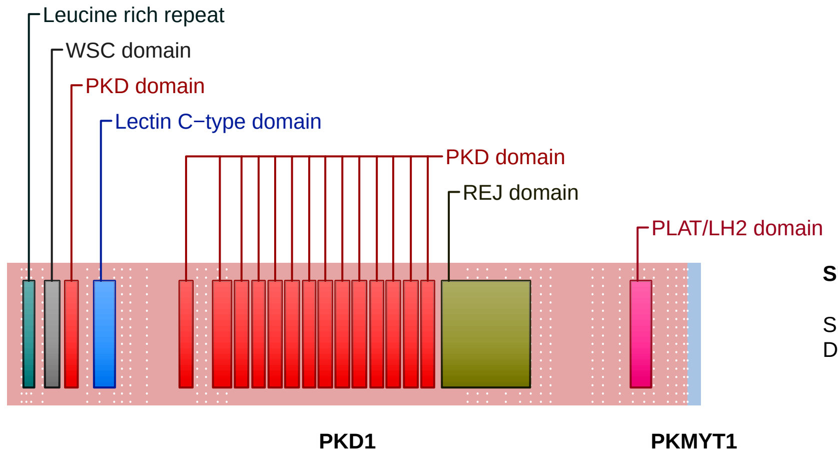


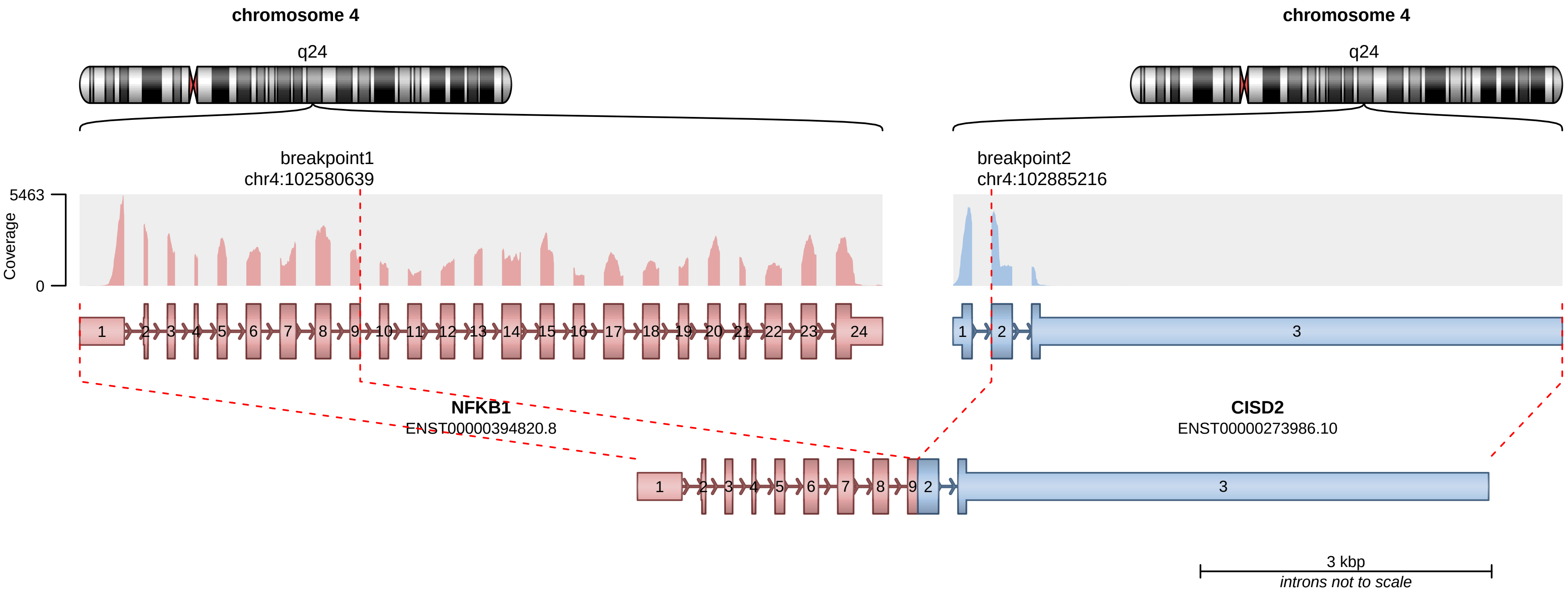


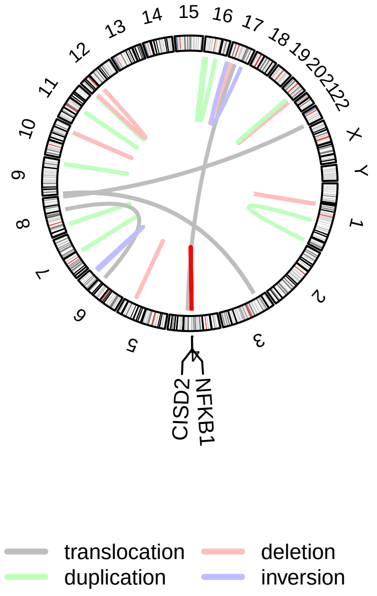
RETAINED PROTEIN DOMAINS
reading frame unclear



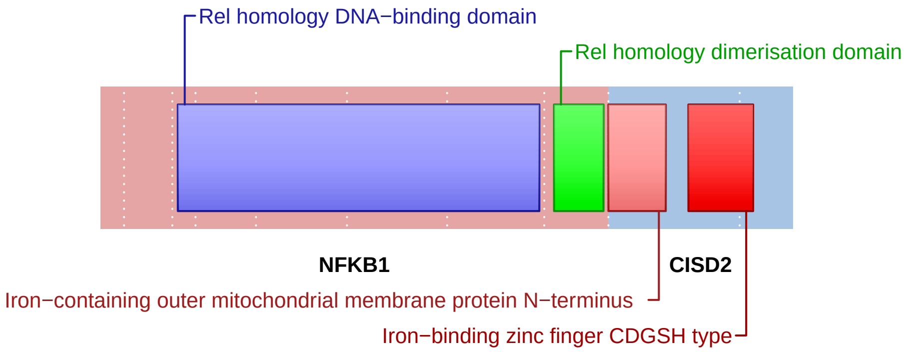
SUPPORTING READ COUNT

Split reads = 299
Discordant mates = 1



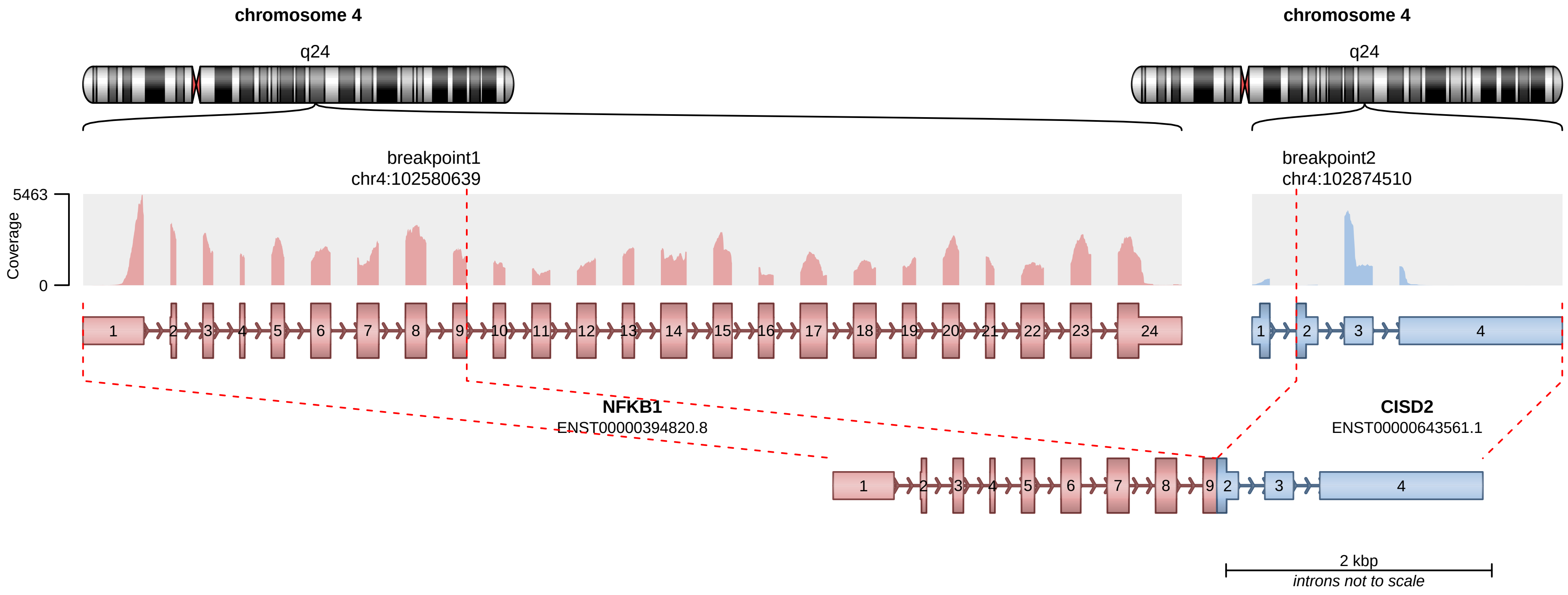


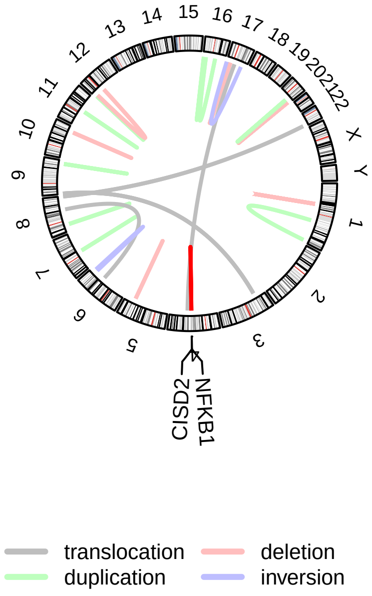
RETAINED PROTEIN DOMAINS
reading frame unclear

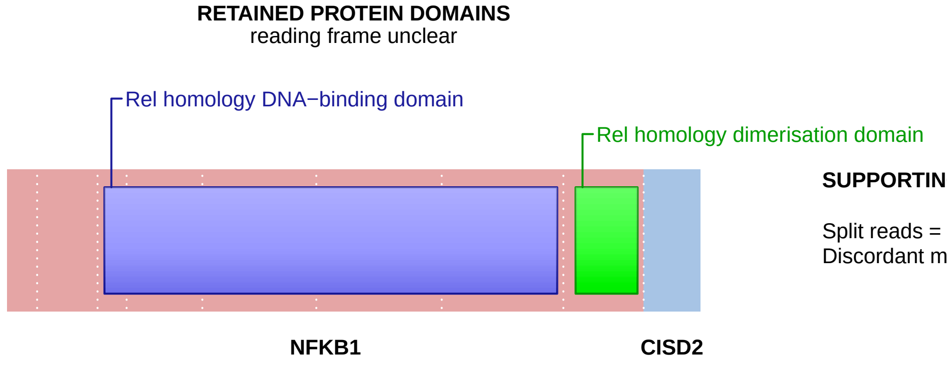


SUPPORTING READ COUNT

Split reads = 227
Discordant mates = 2

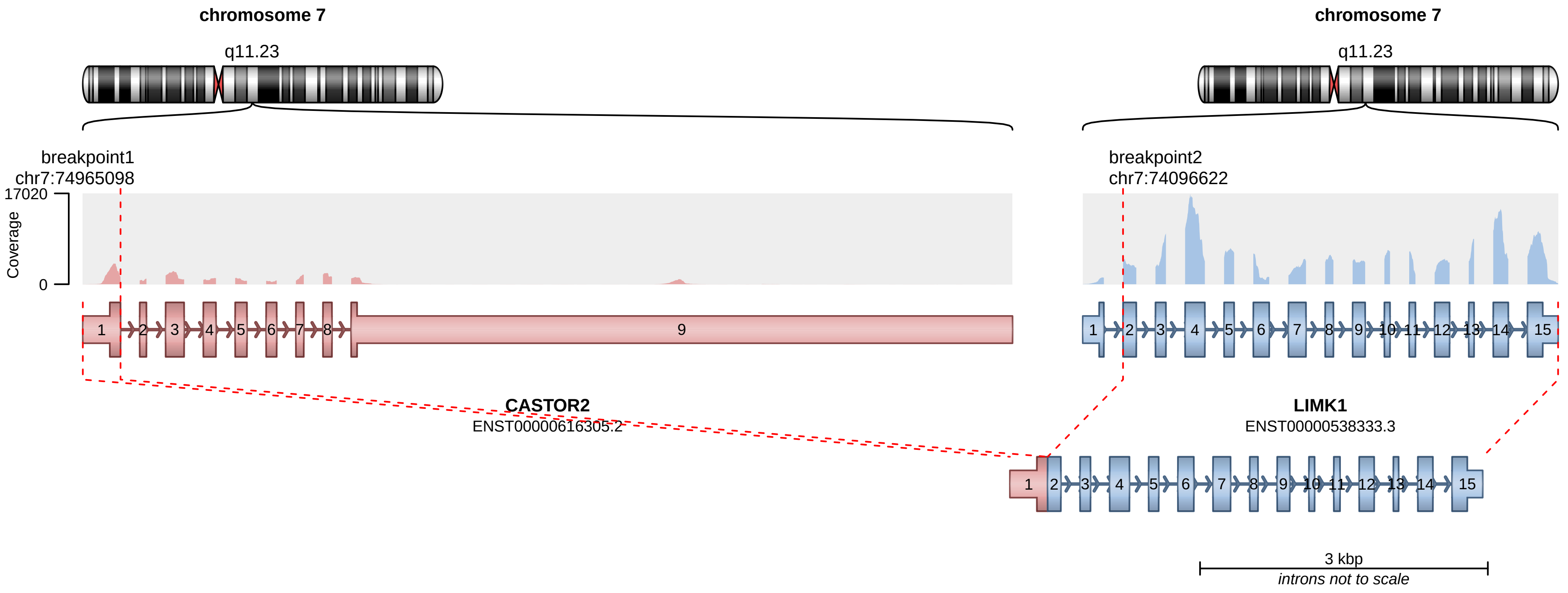


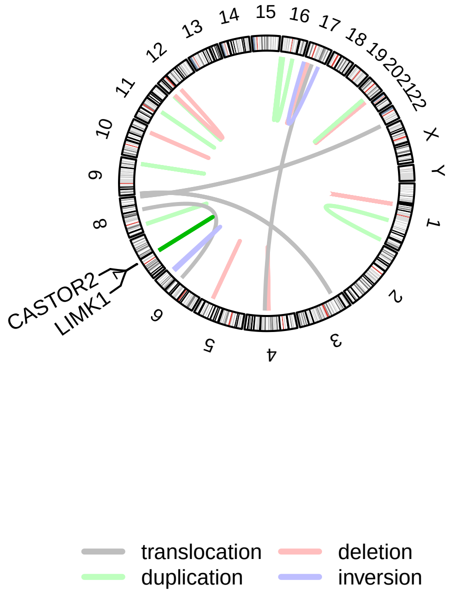


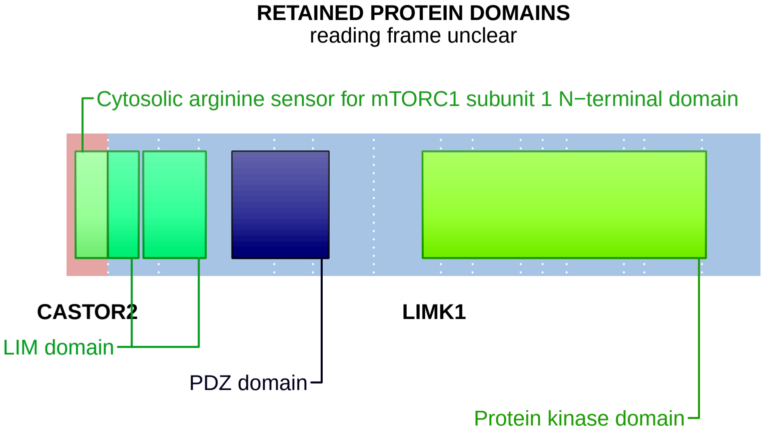


SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 2

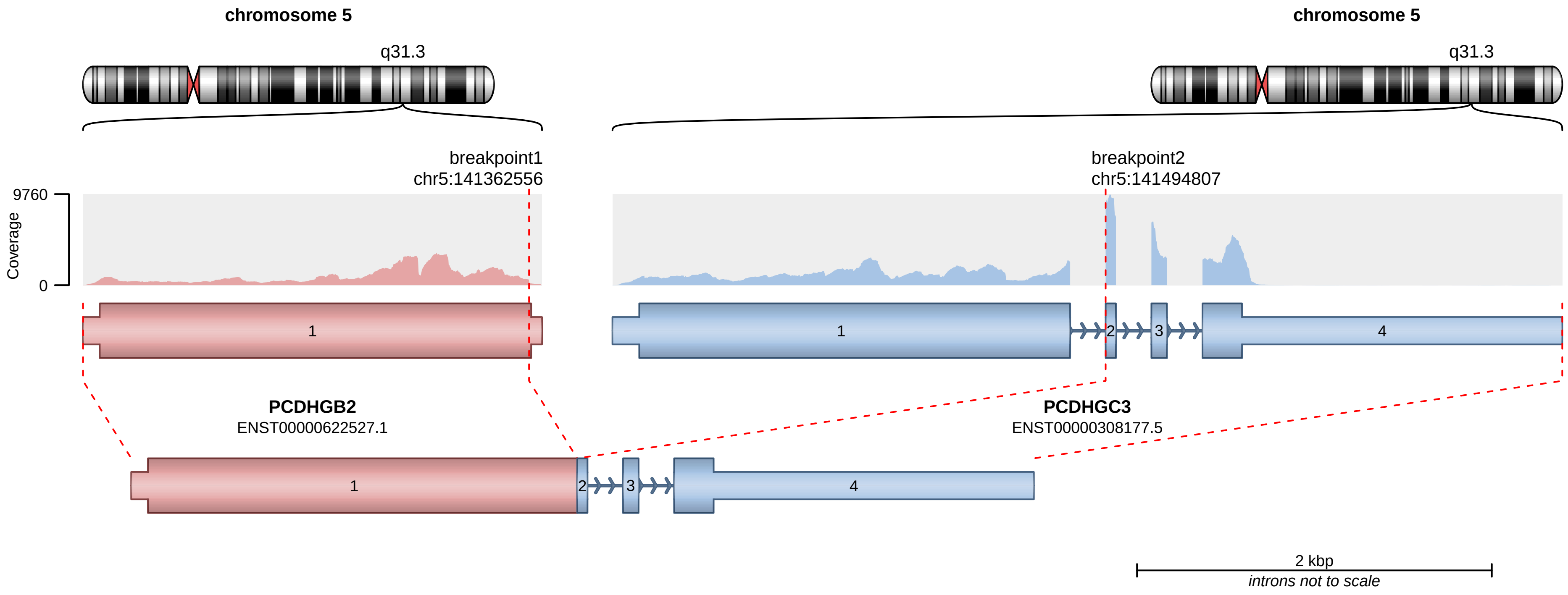


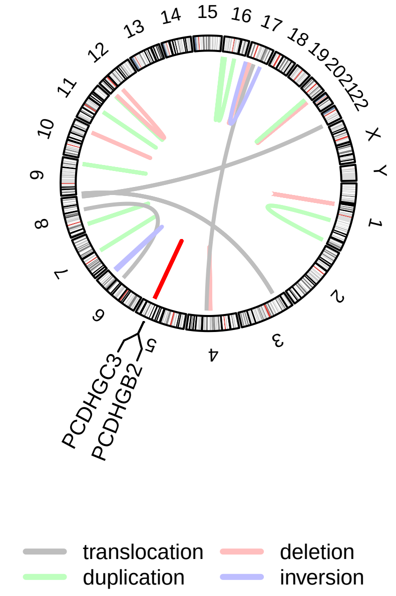




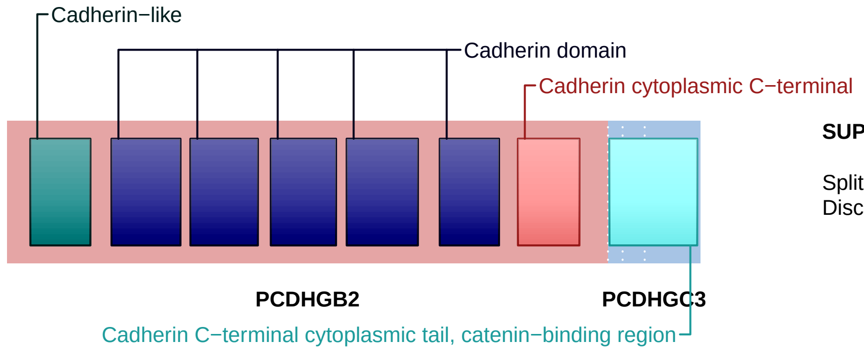
SUPPORTING READ COUNT

Split reads = 150
Discordant mates = 2



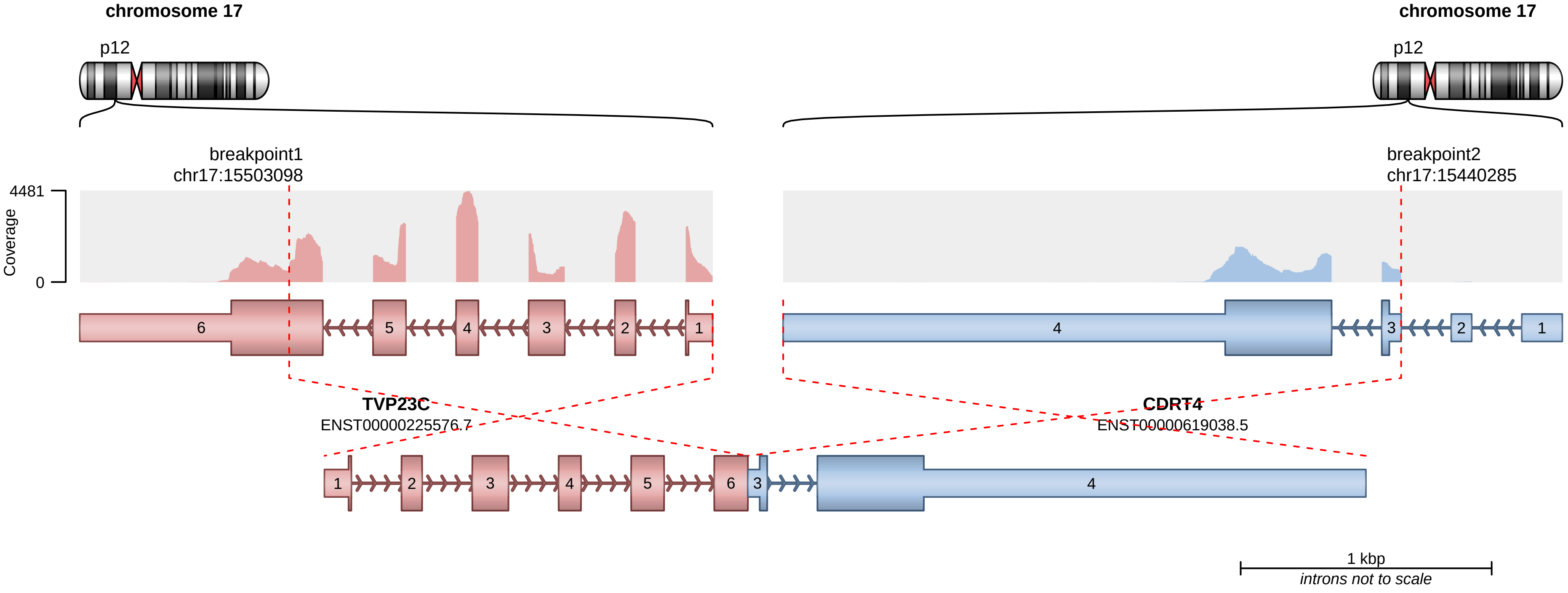


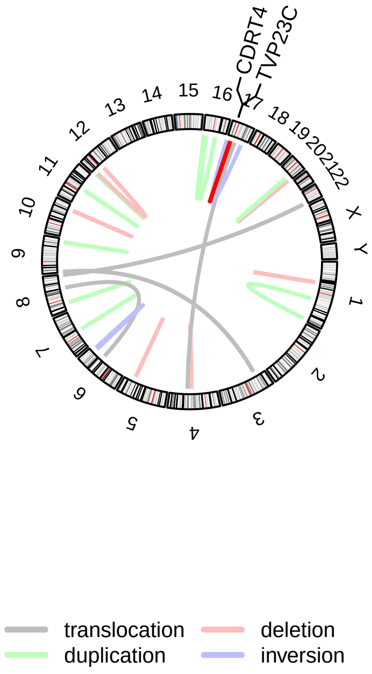
RETAINED PROTEIN DOMAINS
reading frame unclear



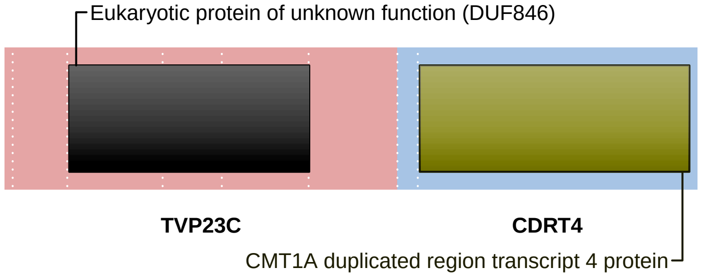
SUPPORTING READ COUNT

Split reads = 126
Discordant mates = 0



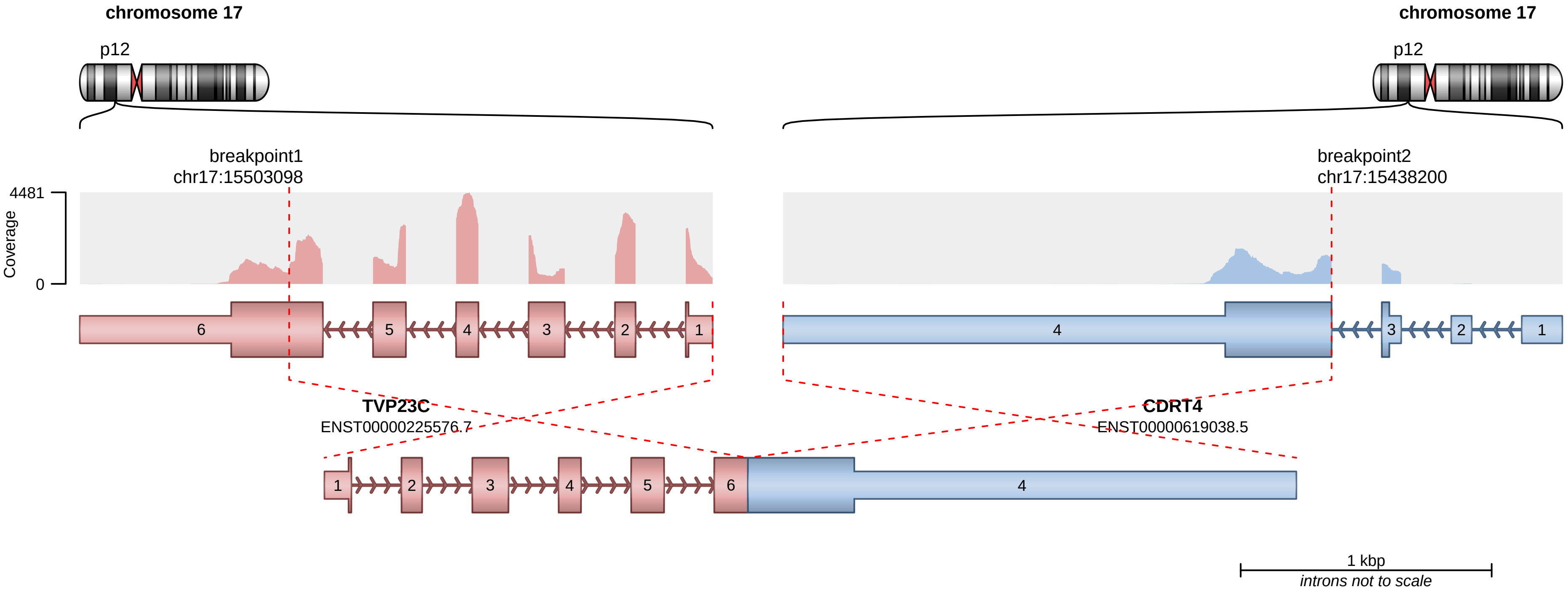


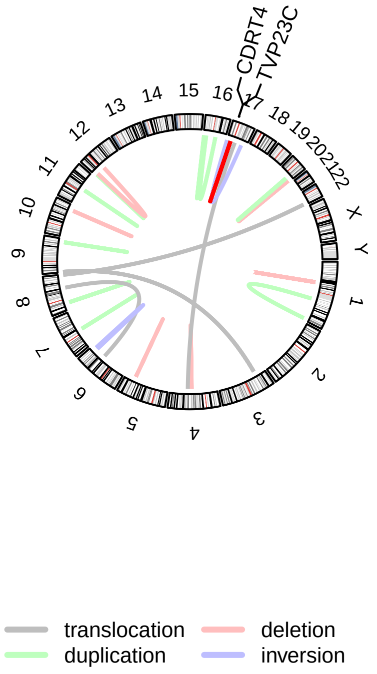
RETAINED PROTEIN DOMAINS
reading frame unclear

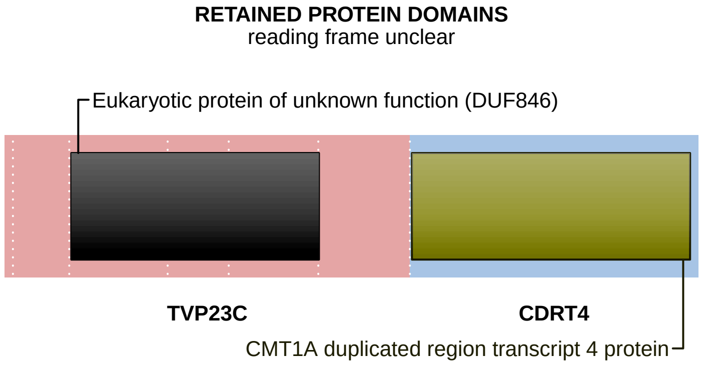


SUPPORTING READ COUNT

Split reads = 119
Discordant mates = 1

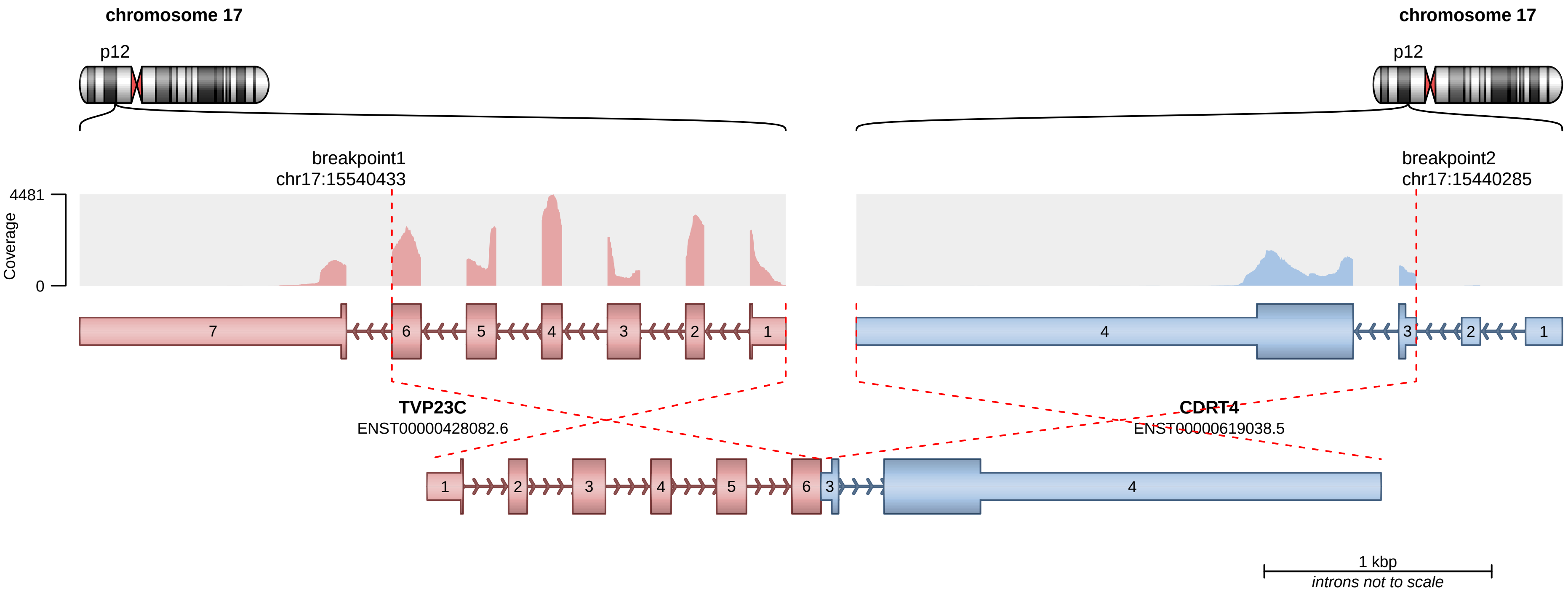


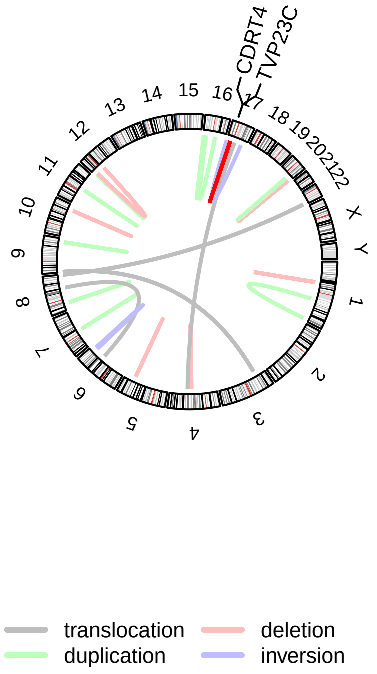




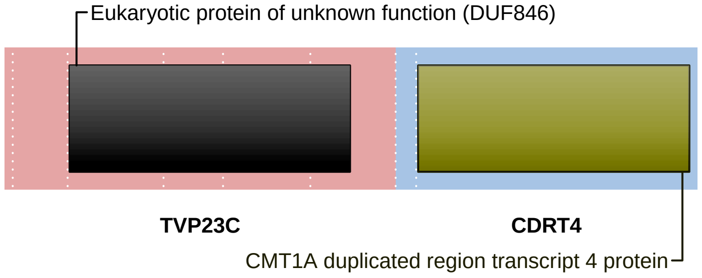
SUPPORTING READ COUNT

Split reads = 44
Discordant mates = 0



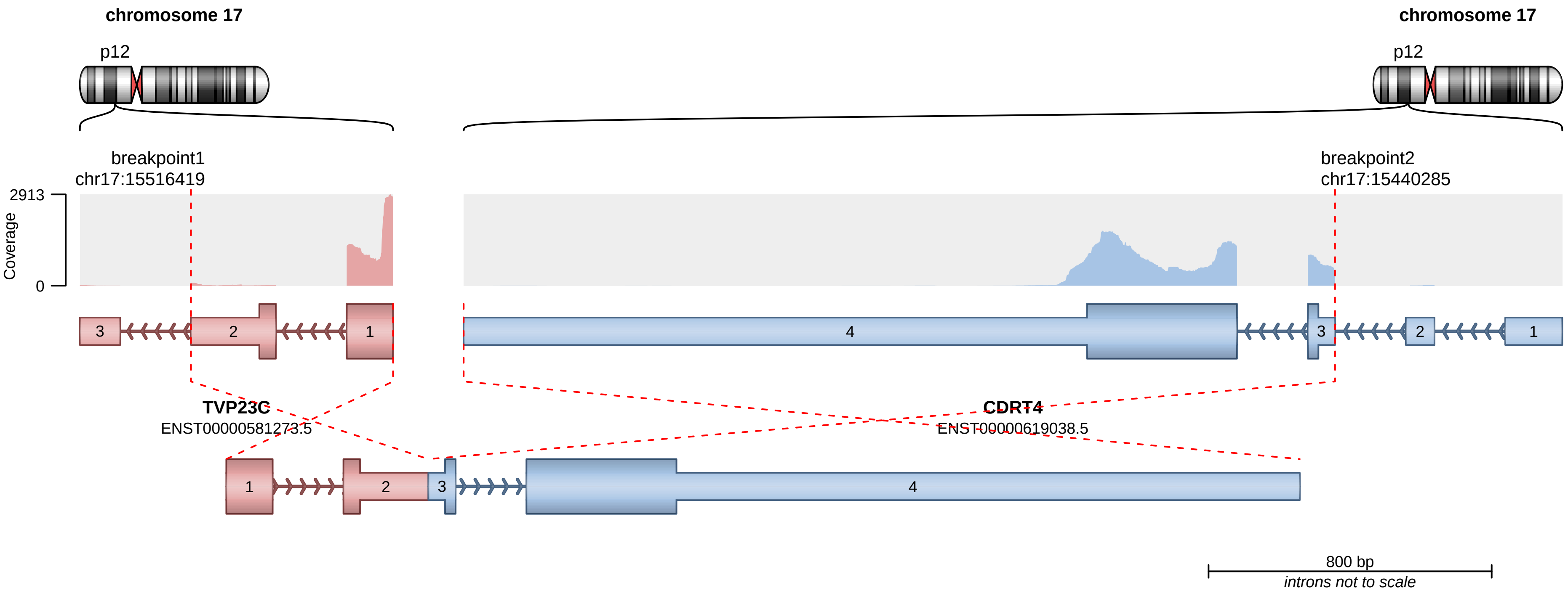


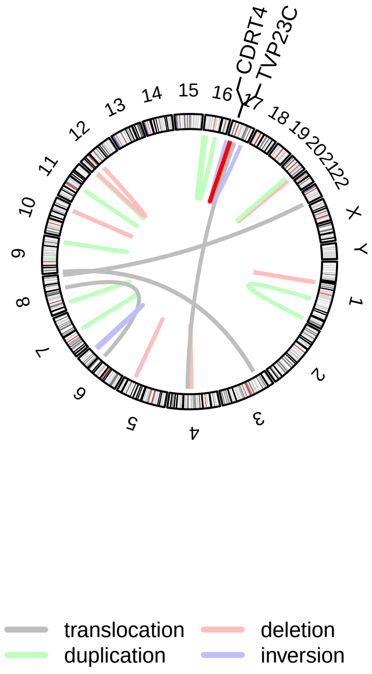
RETAINED PROTEIN DOMAINS
reading frame unclear



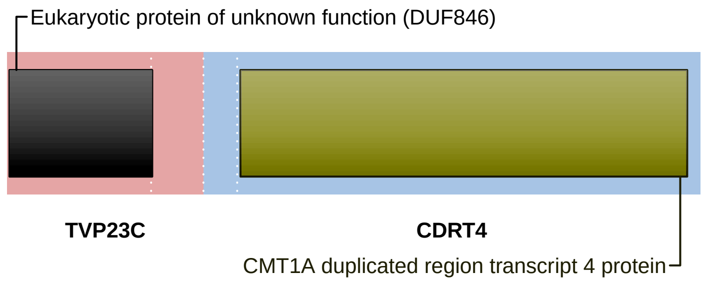
SUPPORTING READ COUNT

Split reads = 36
Discordant mates = 0



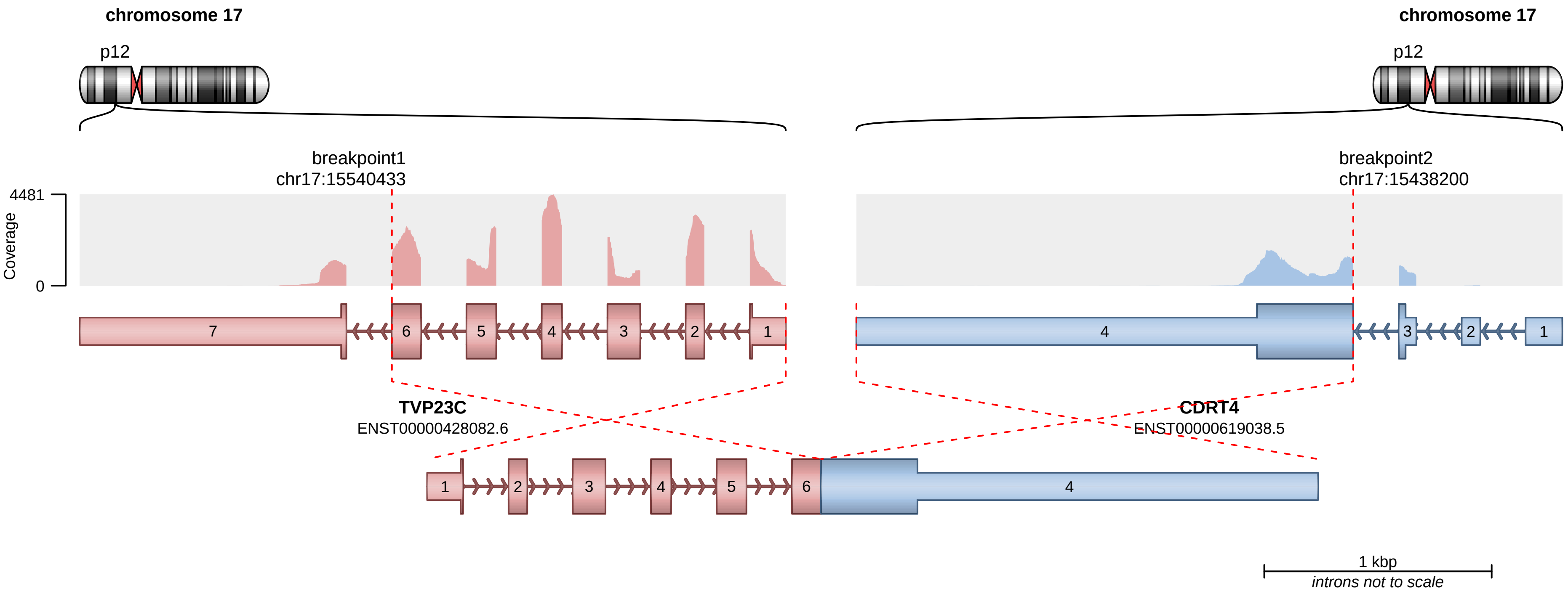


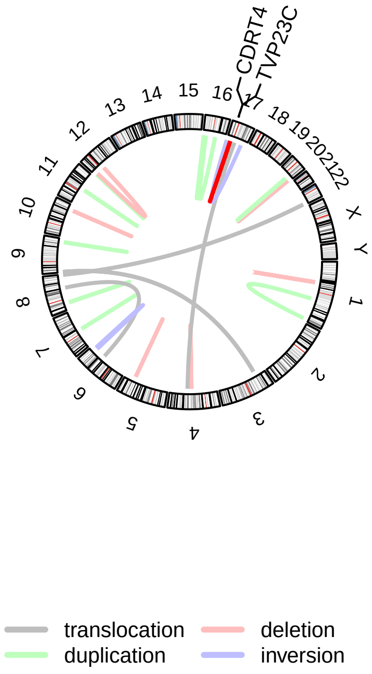
RETAINED PROTEIN DOMAINS
reading frame unclear

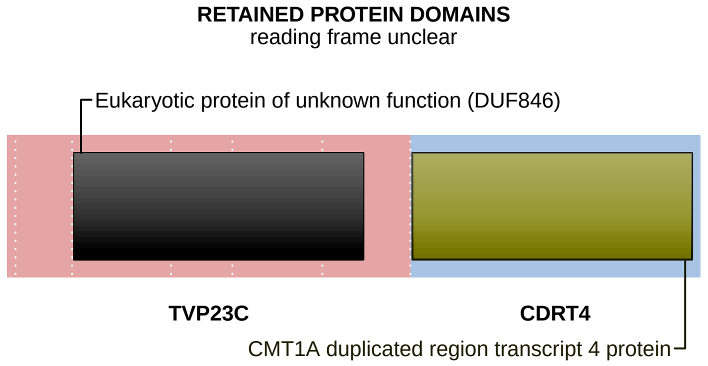


SUPPORTING READ COUNT

Split reads = 21
Discordant mates = 0

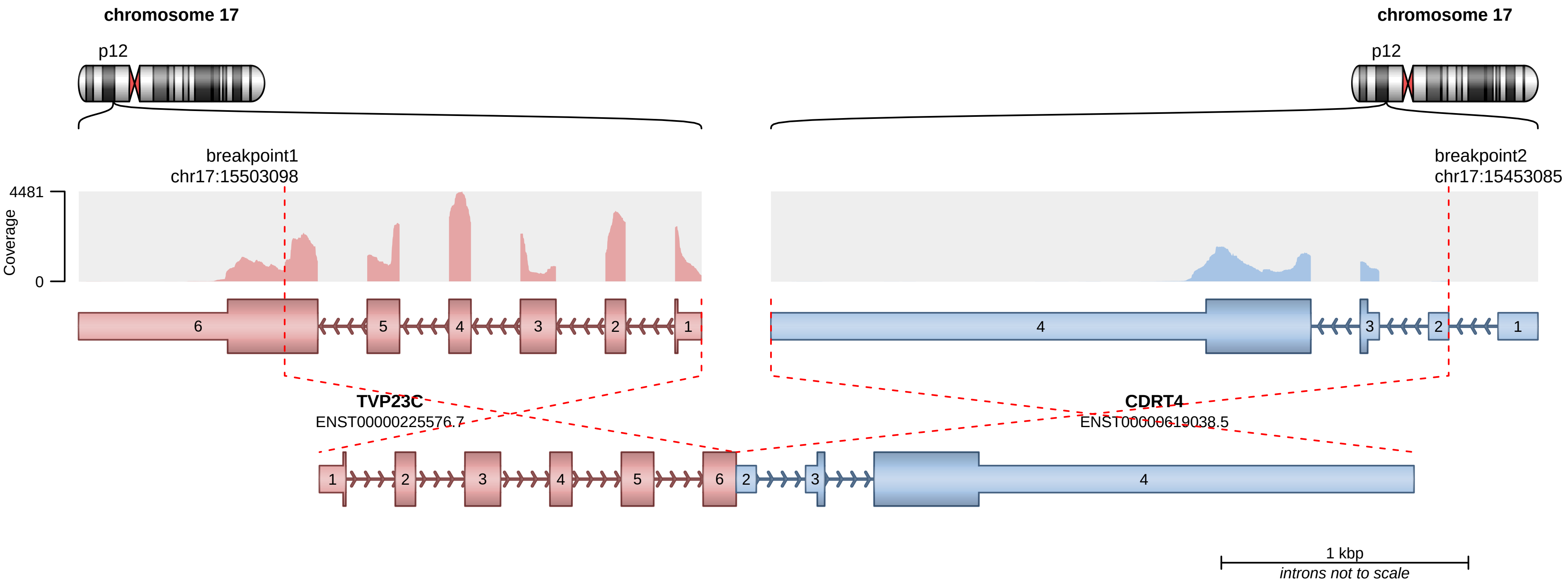


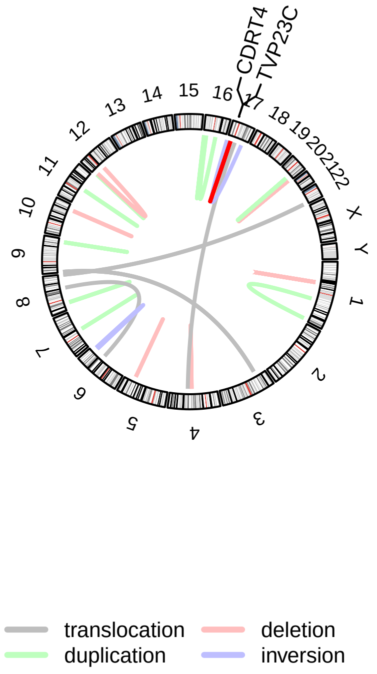


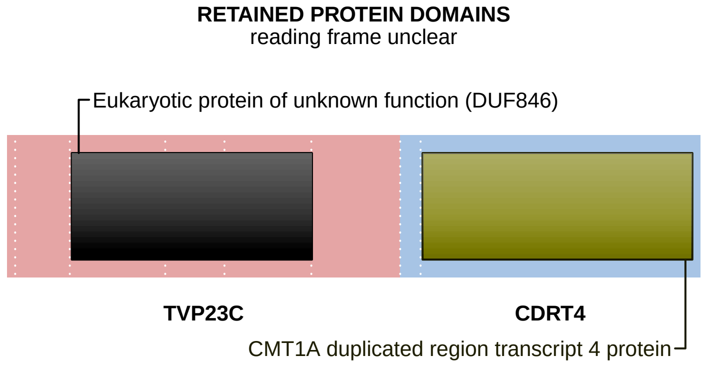


SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

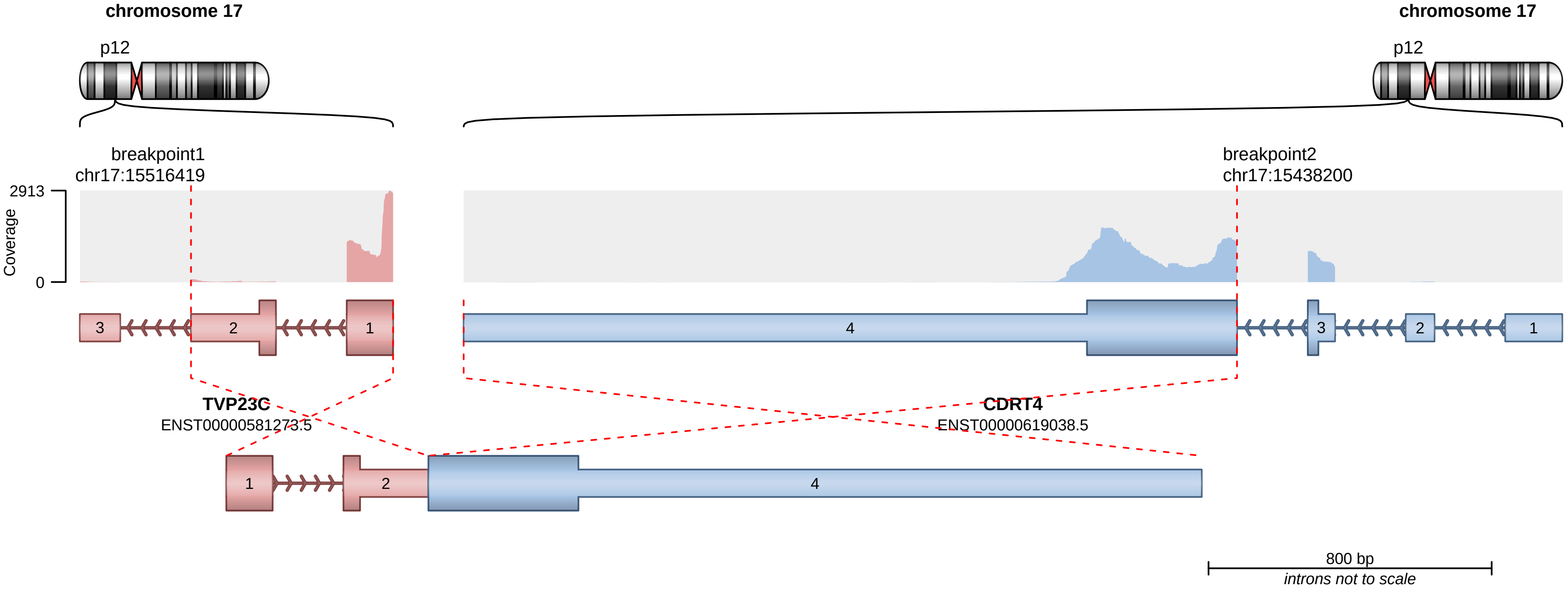


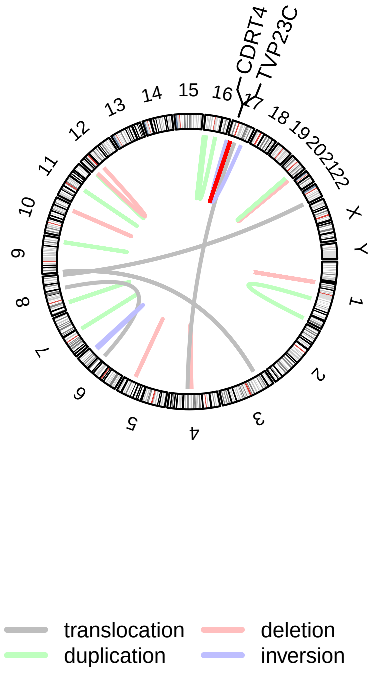


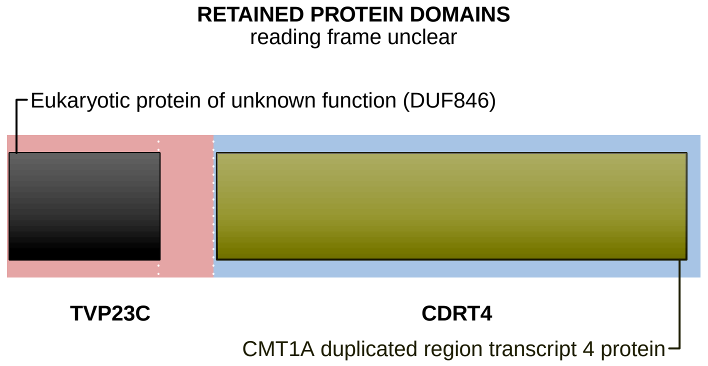


SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 1

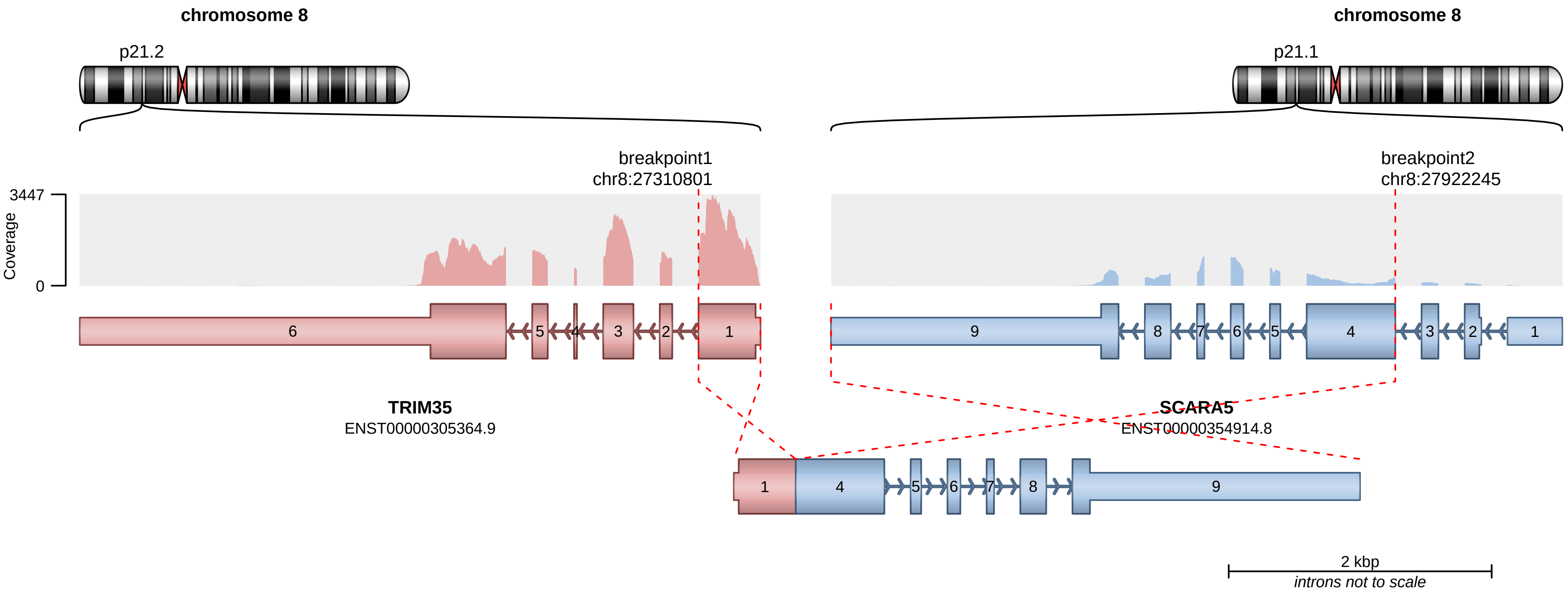


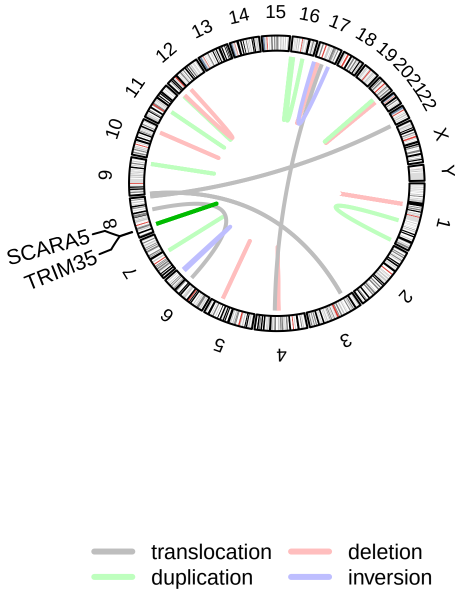


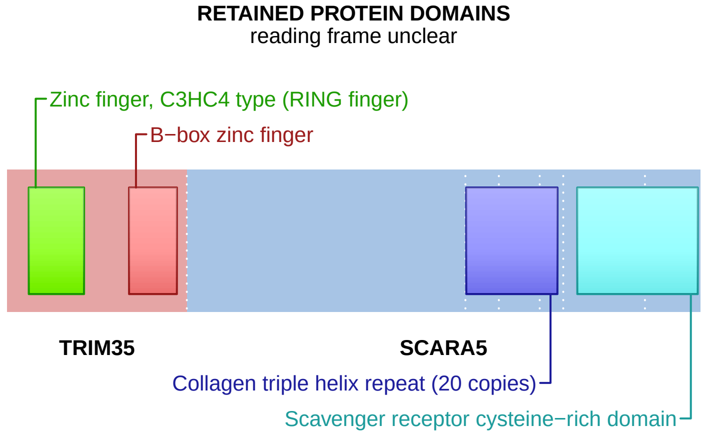


SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

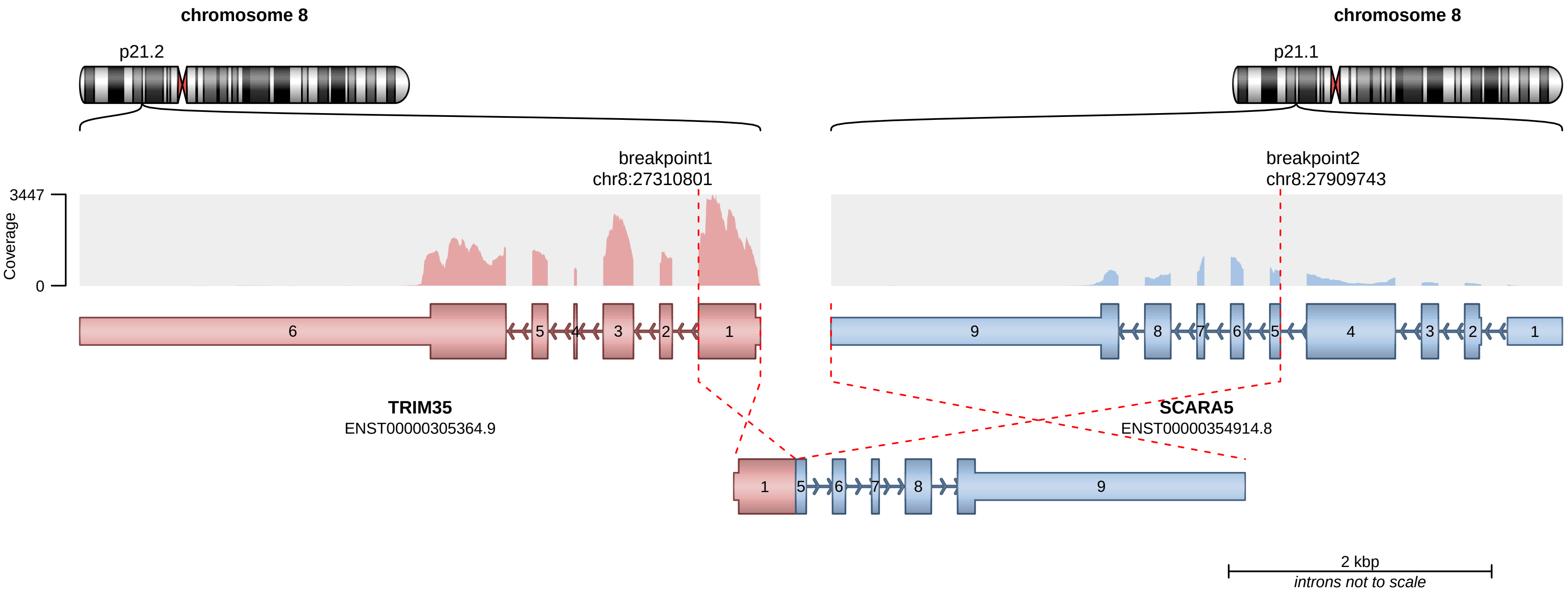


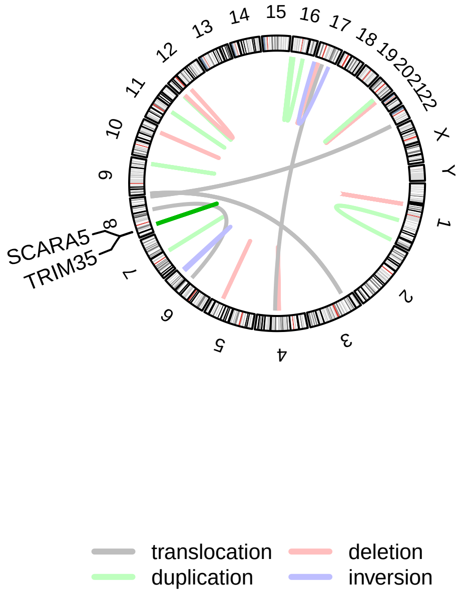




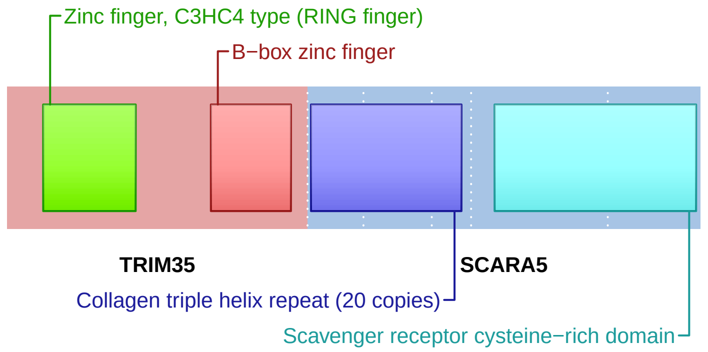
SUPPORTING READ COUNT

Split reads = 114
Discordant mates = 0



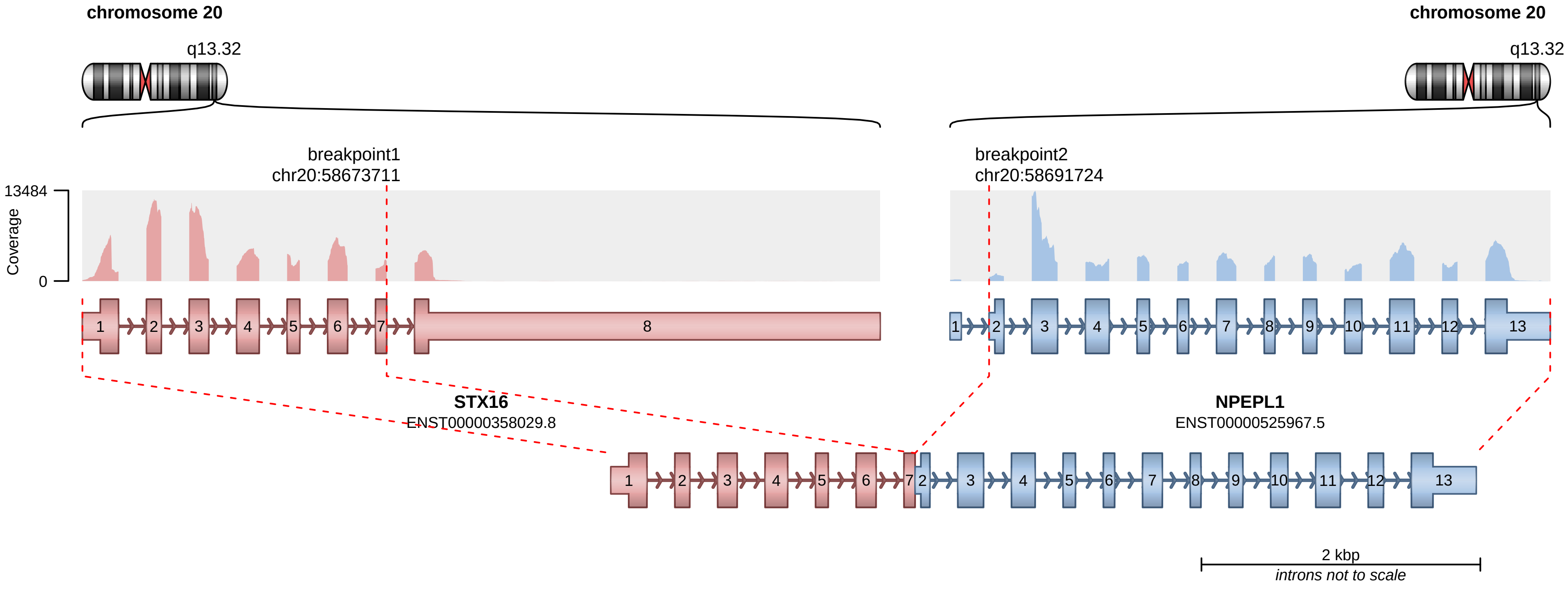


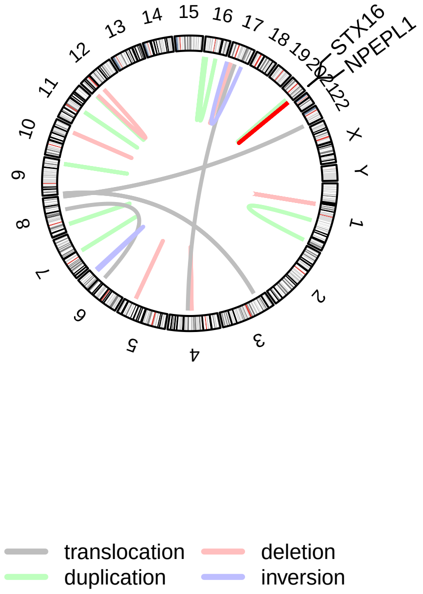
RETAINED PROTEIN DOMAINS
reading frame unclear

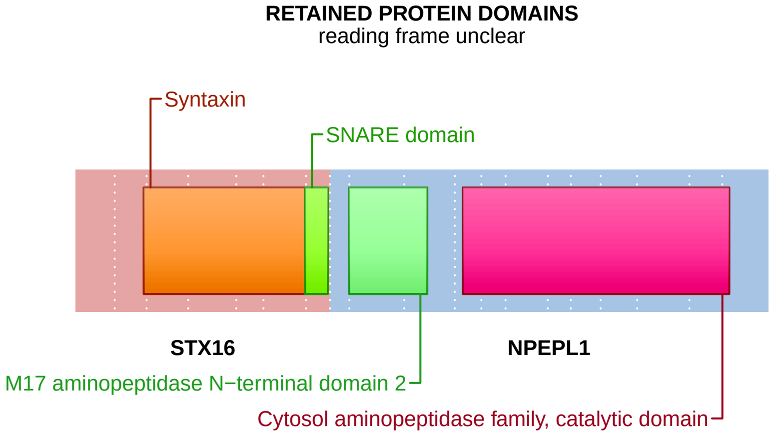


SUPPORTING READ COUNT

Split reads = 4
Discordant mates = 0

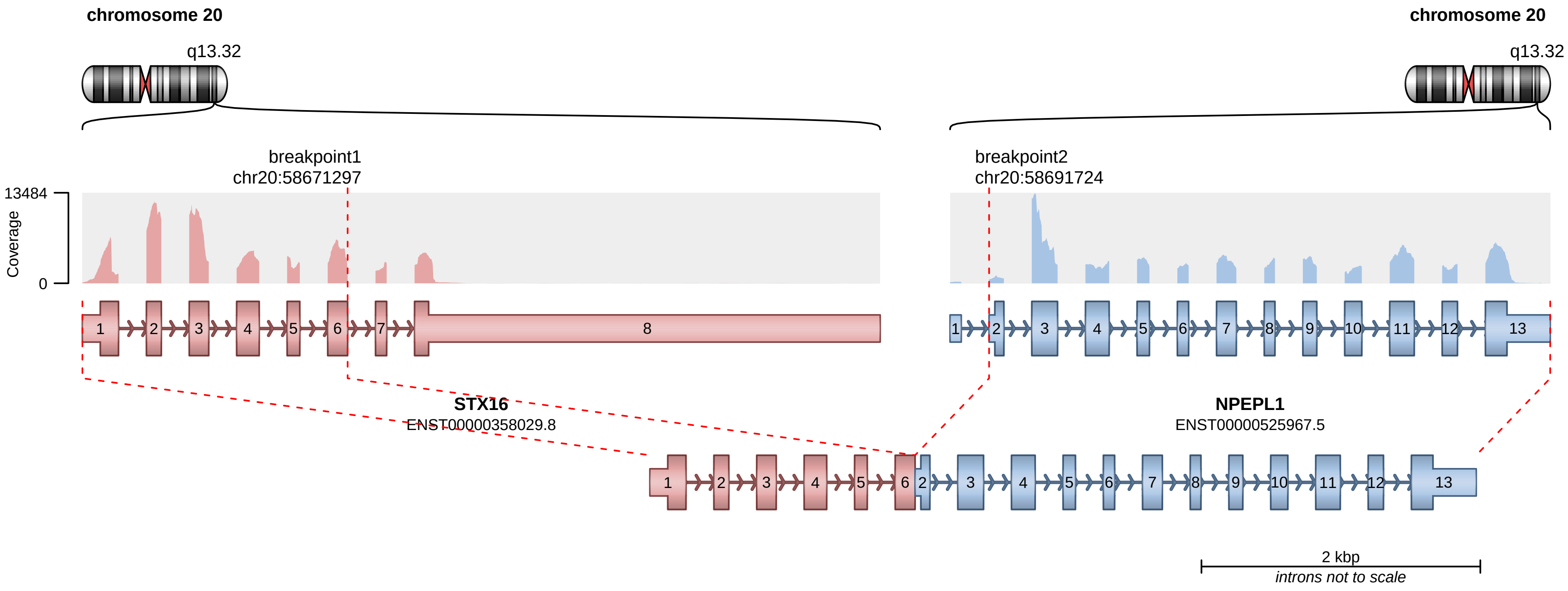


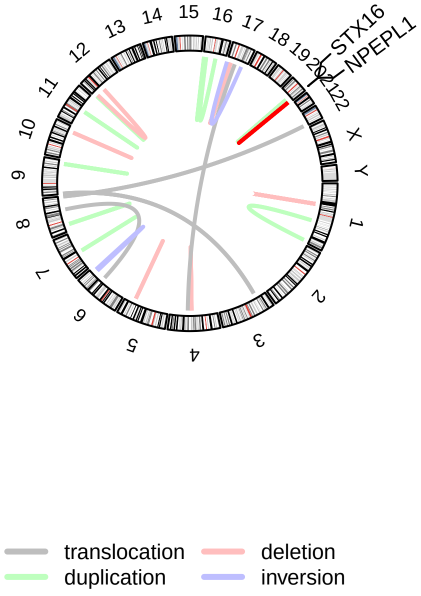


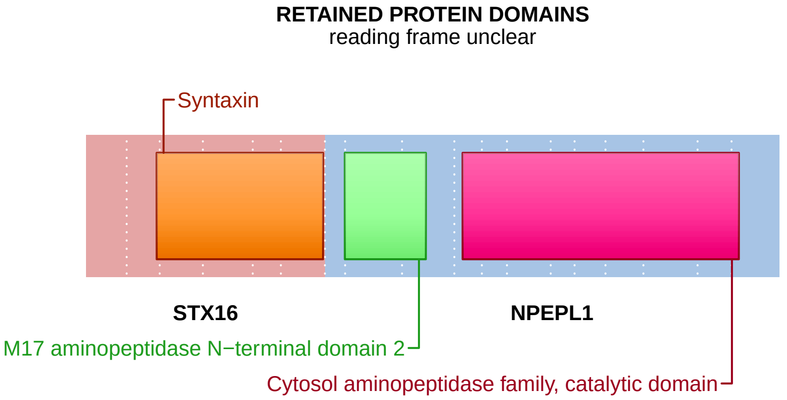


SUPPORTING READ COUNT

Split reads = 111
Discordant mates = 2

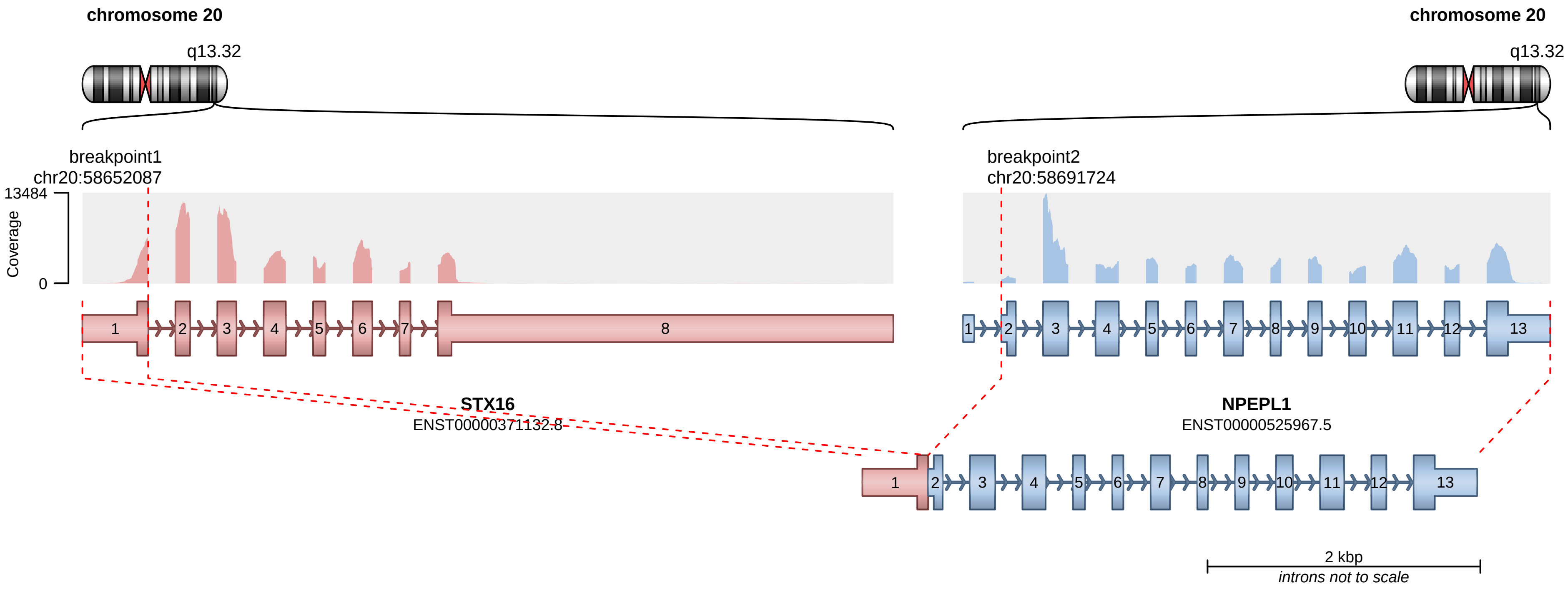


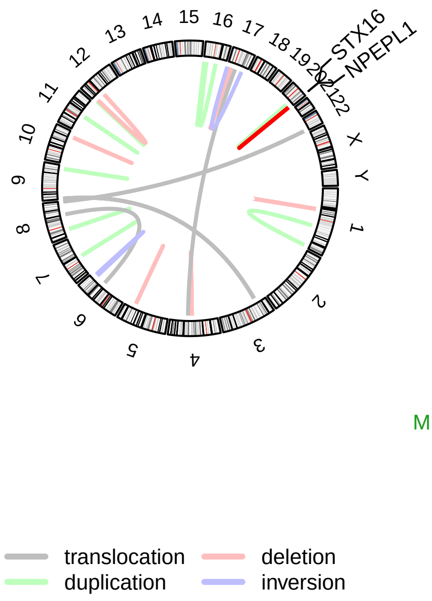


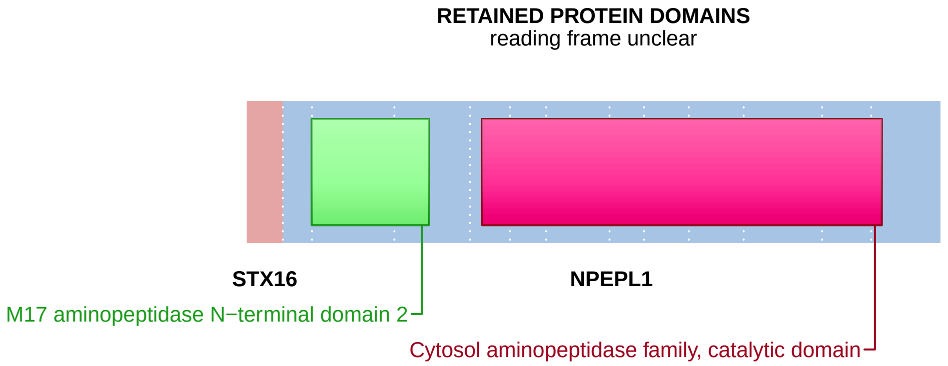


SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

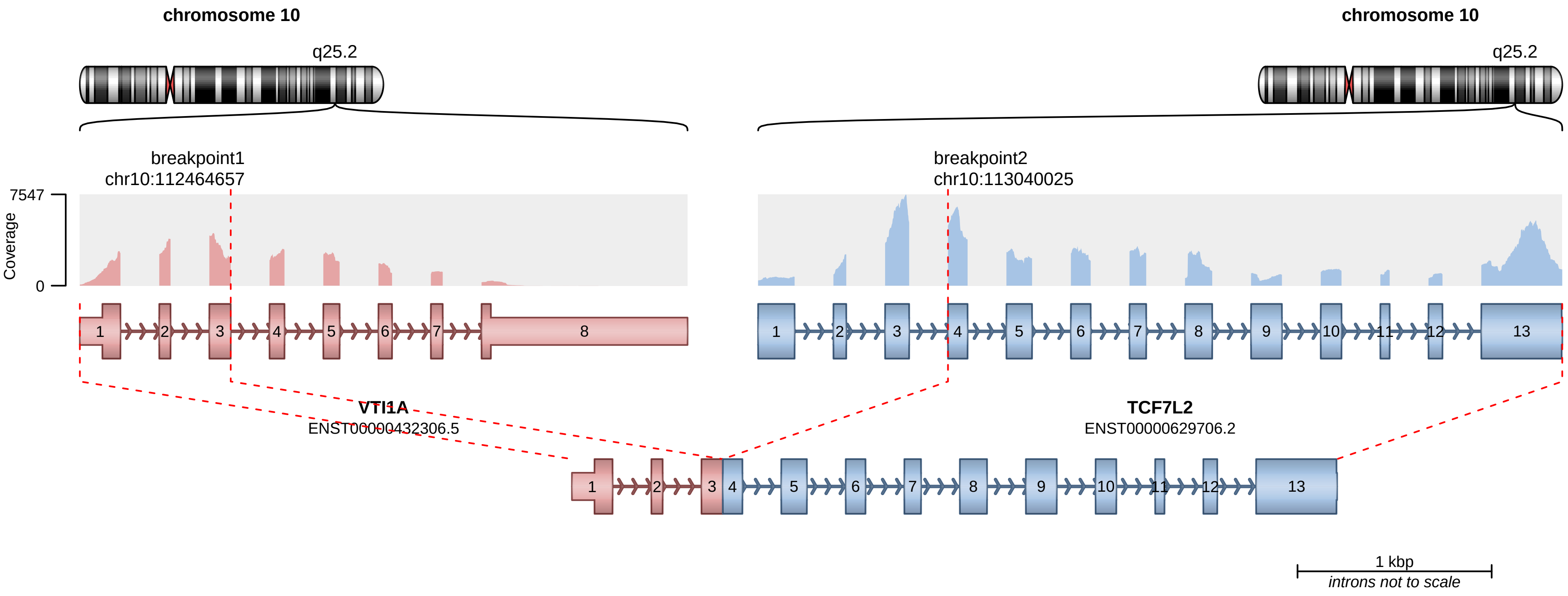


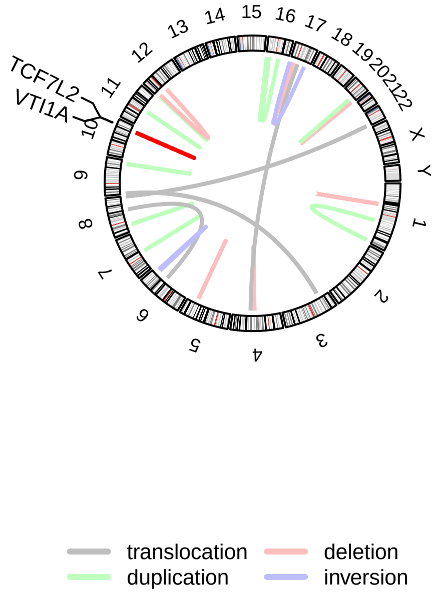


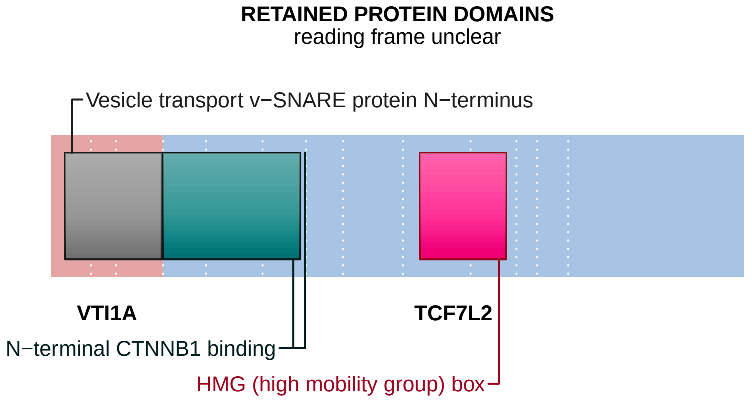


SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

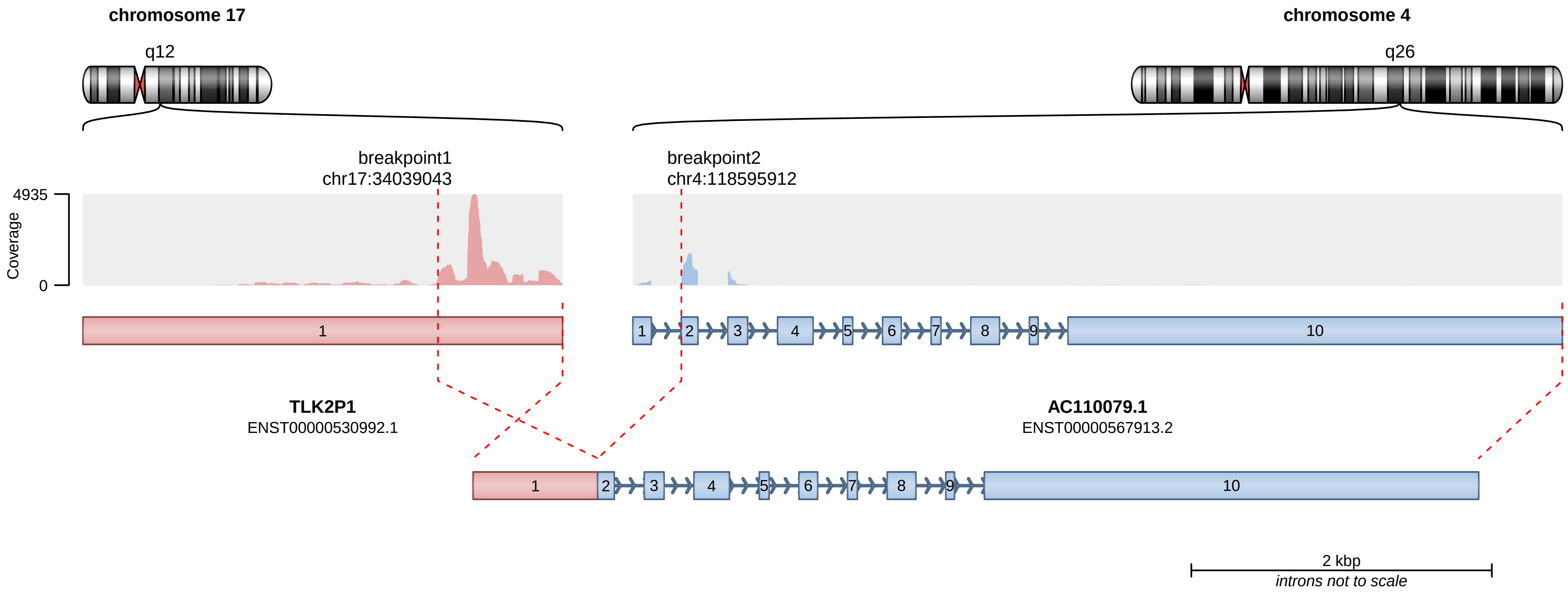


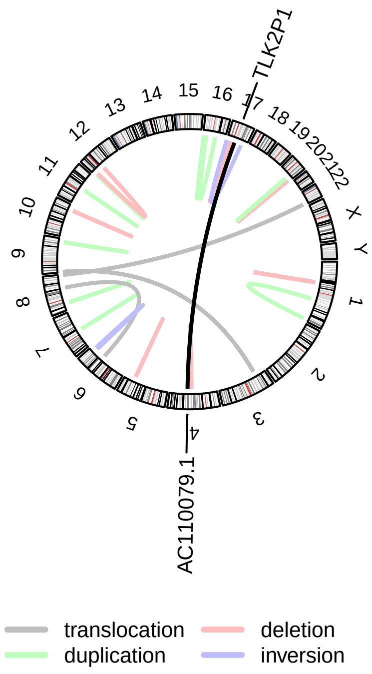




SUPPORTING READ COUNT

Split reads = 98
Discordant mates = 0

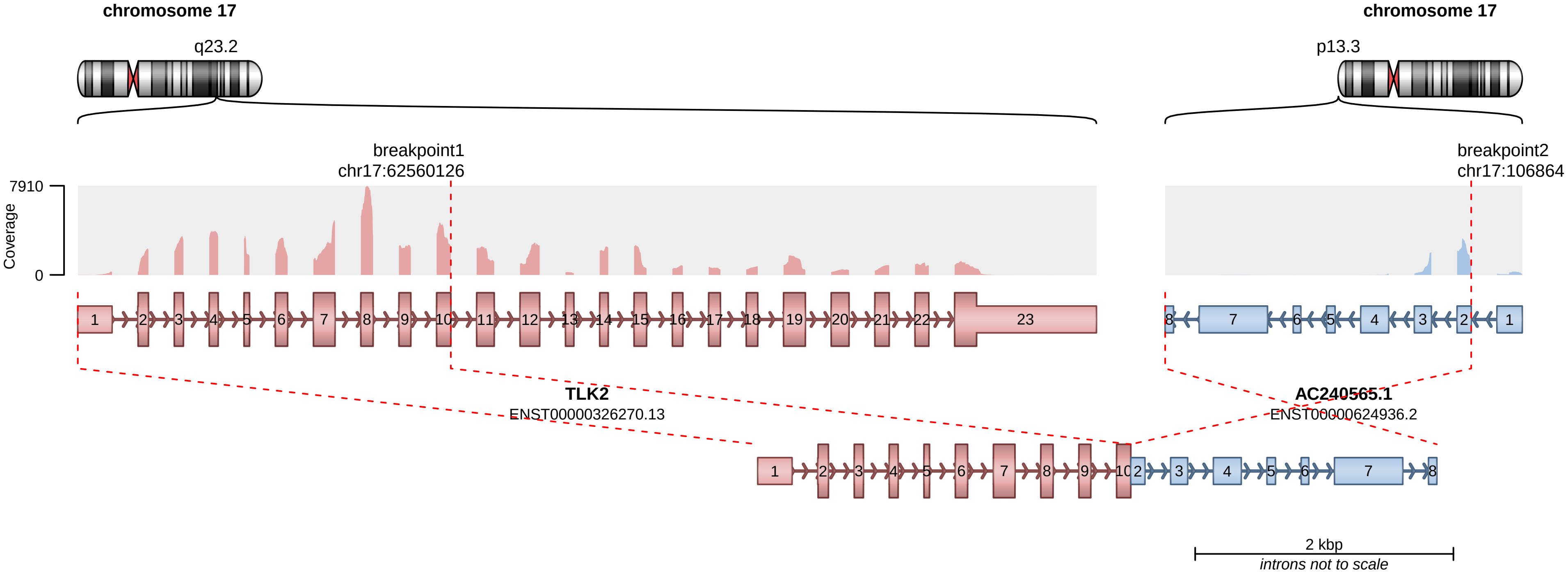


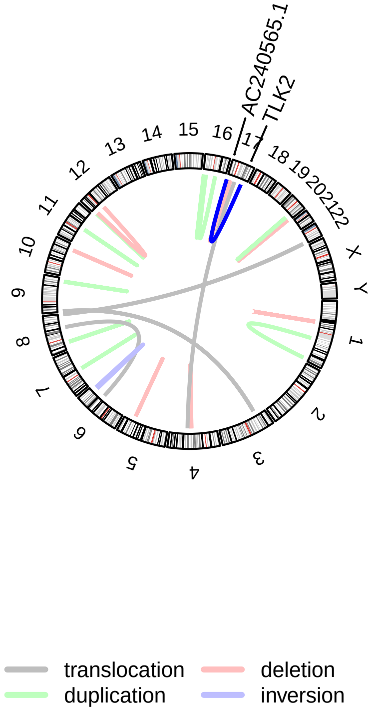


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 91
Discordant mates = 0

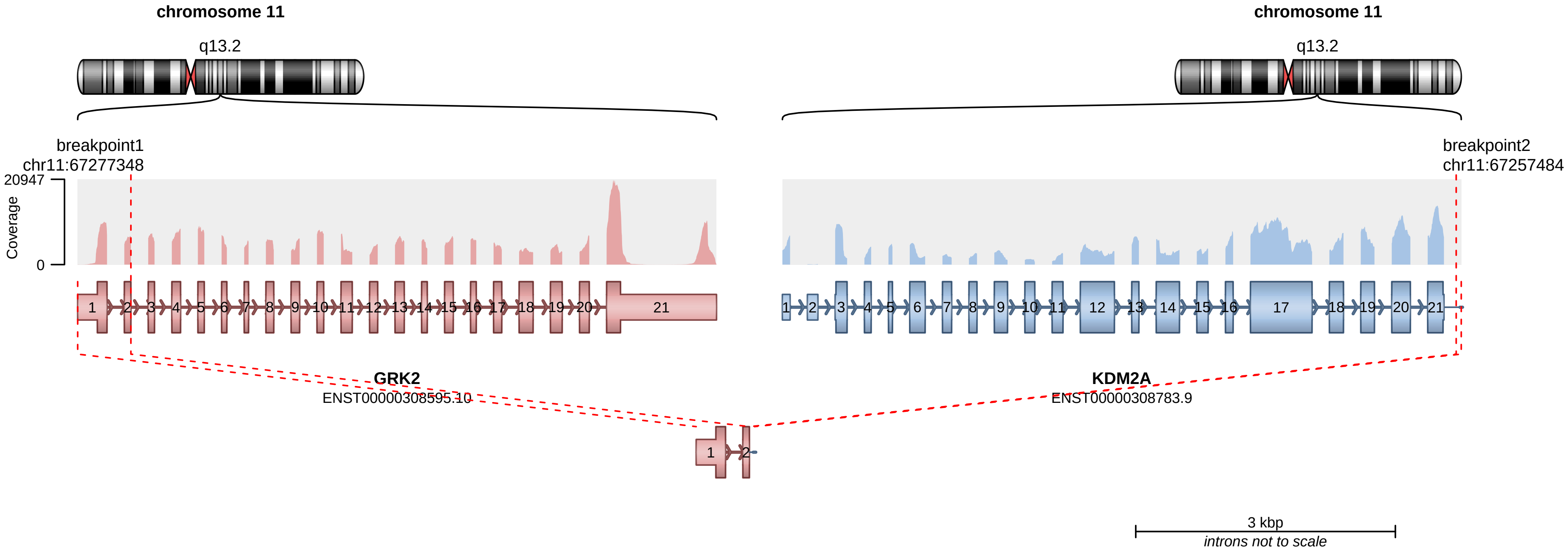


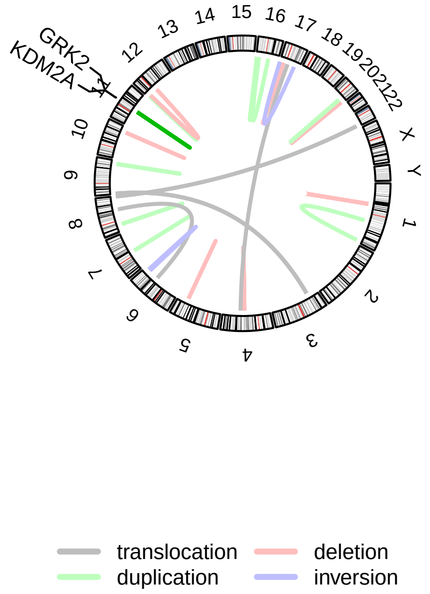


No protein domains retained in fusion.

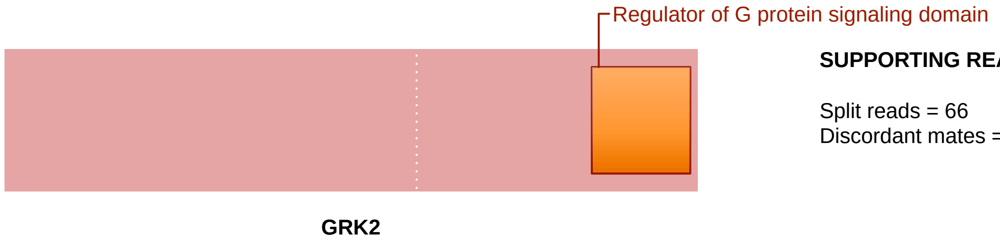
SUPPORTING READ COUNT

Split reads = 85
Discordant mates = 0



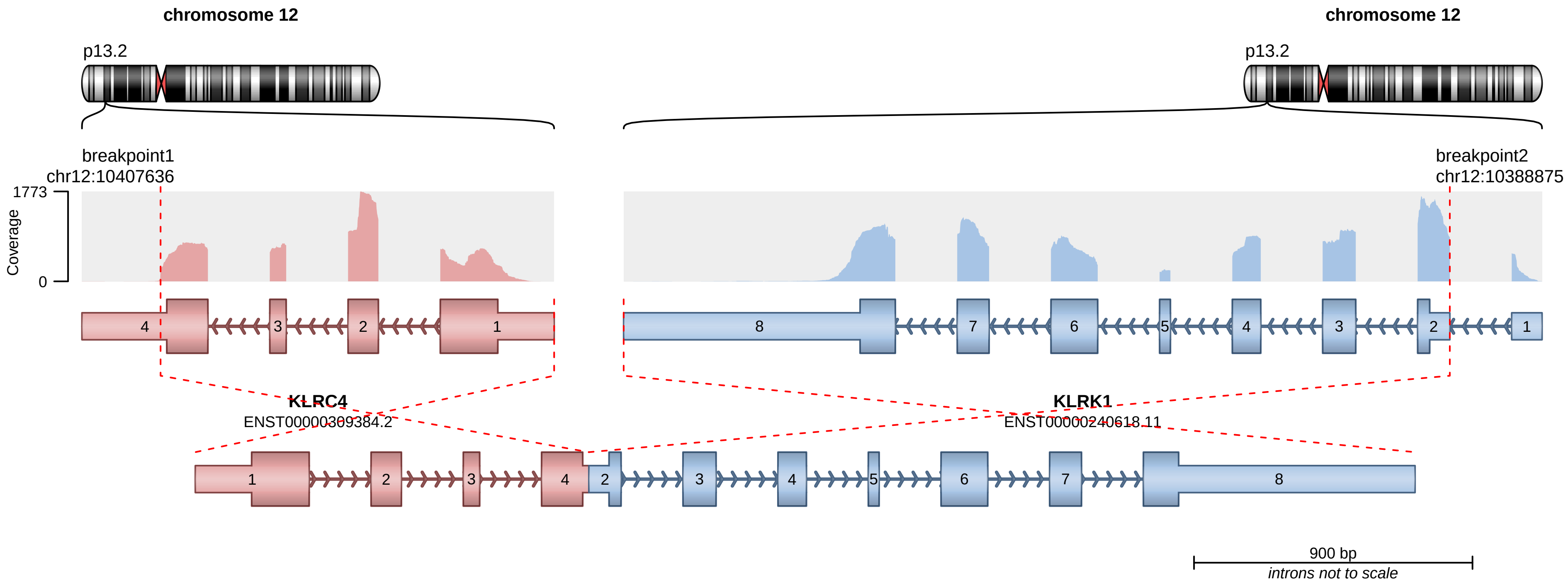


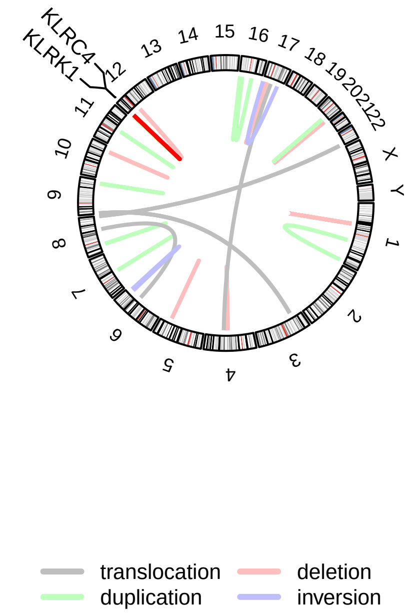
RETAINED PROTEIN DOMAINS
reading frame unclear



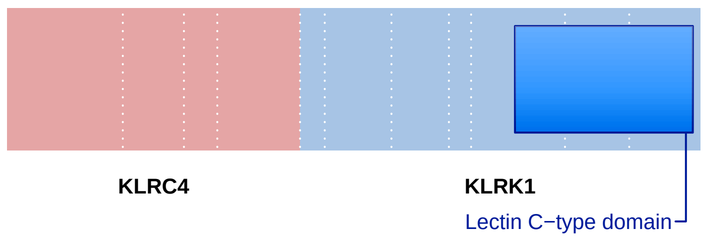
SUPPORTING READ COUNT

Split reads = 66
Discordant mates = 0



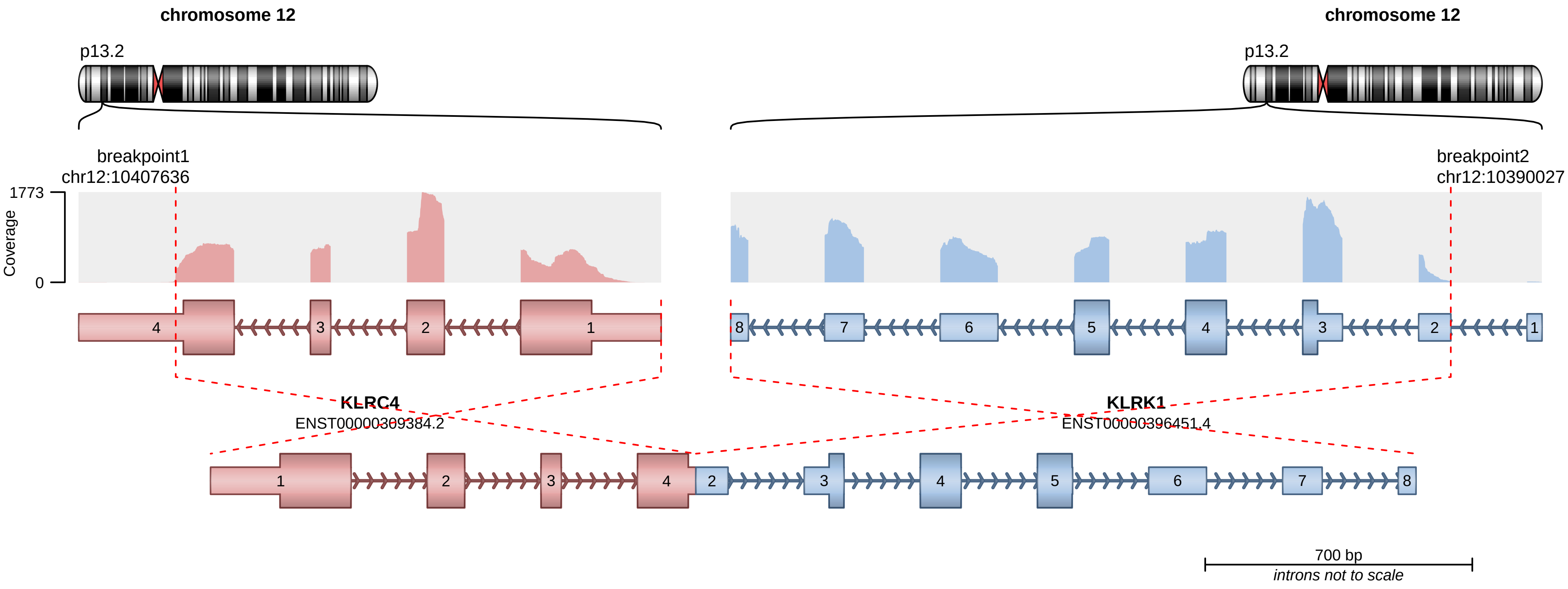


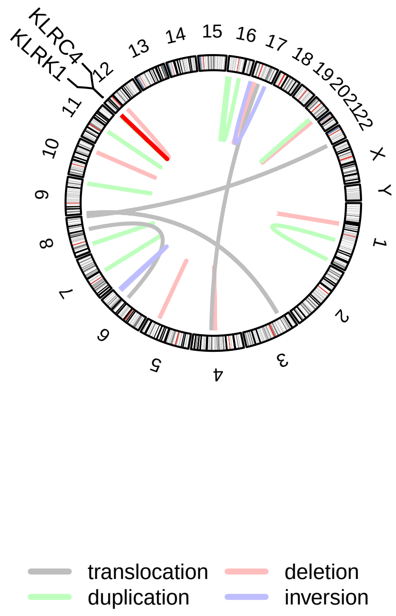
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 51
Discordant mates = 0

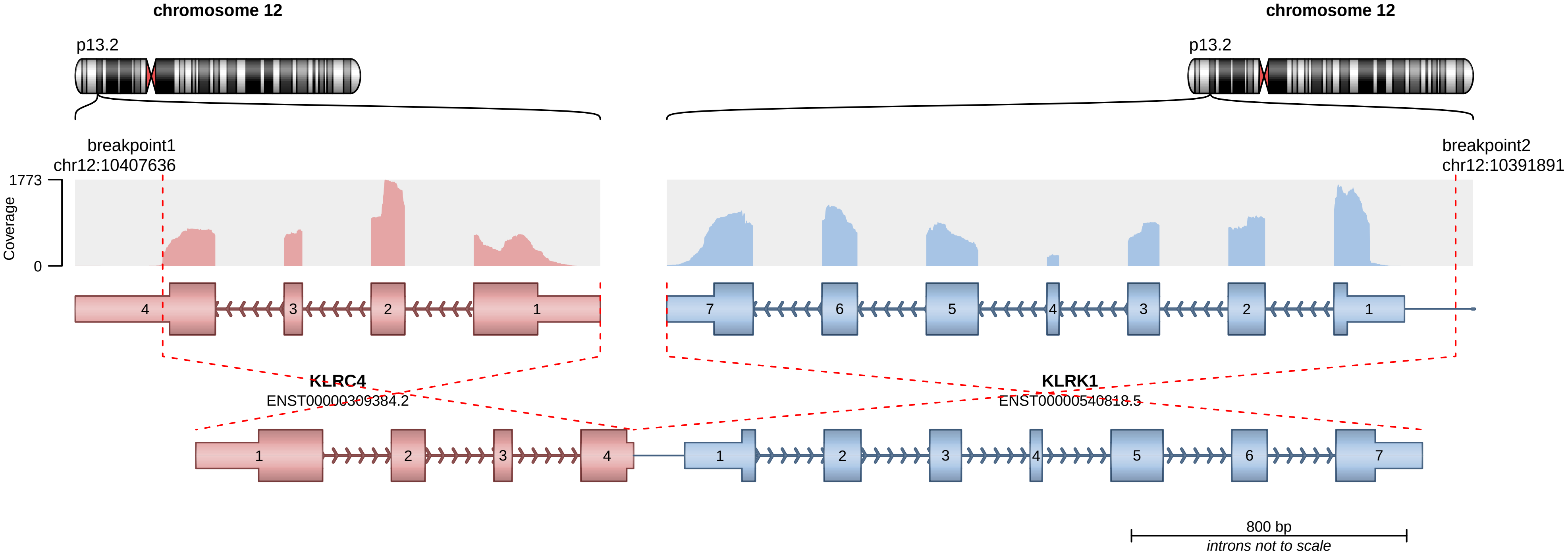


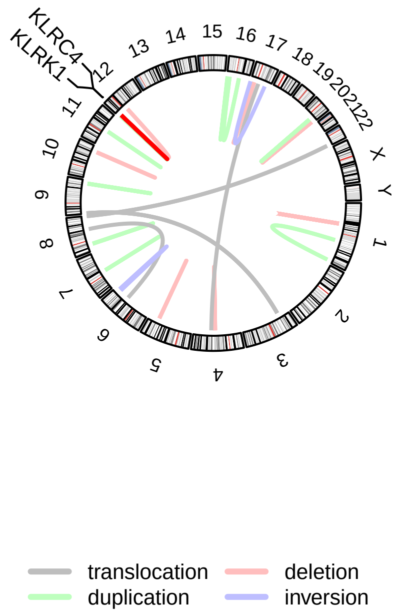


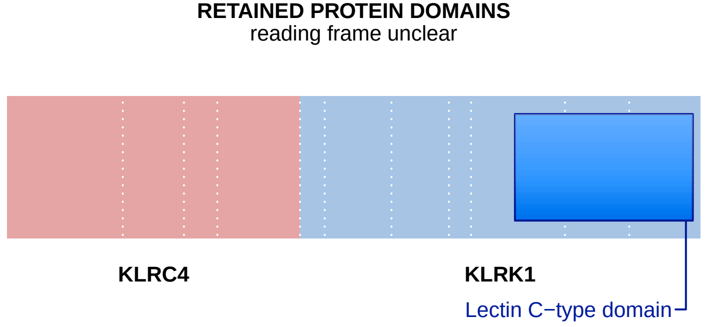
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0

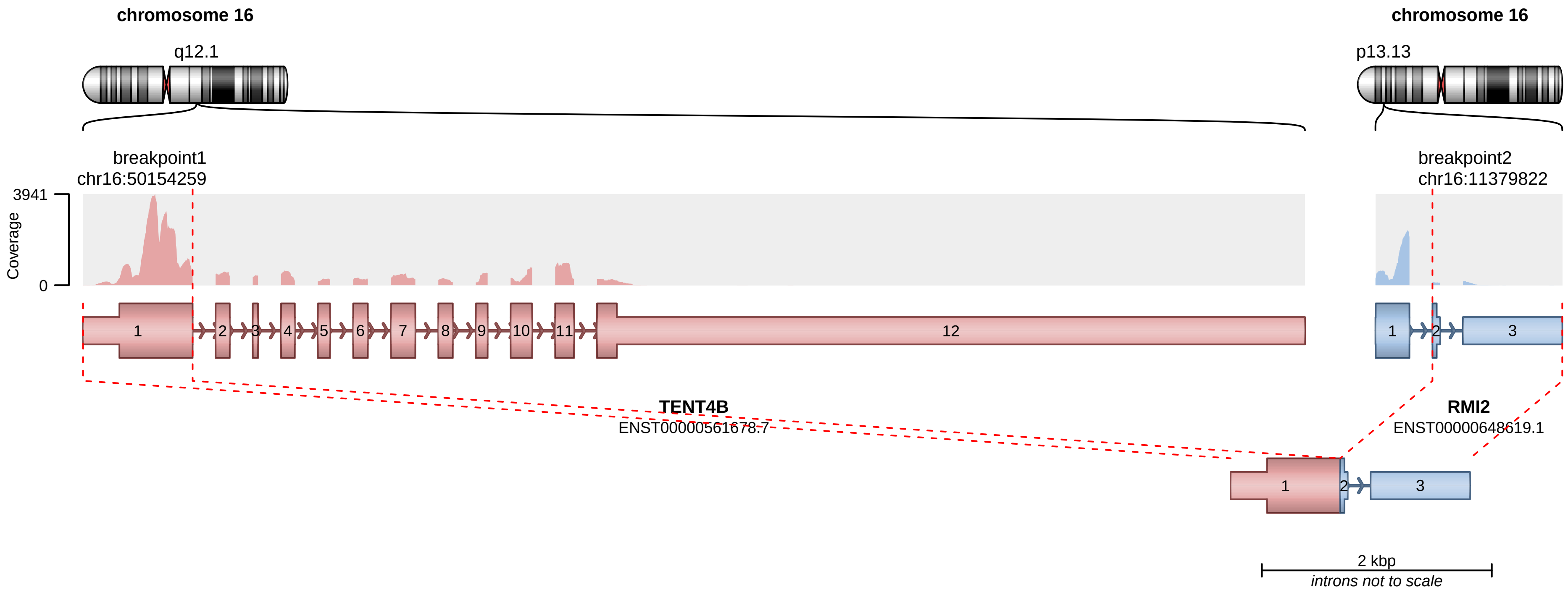


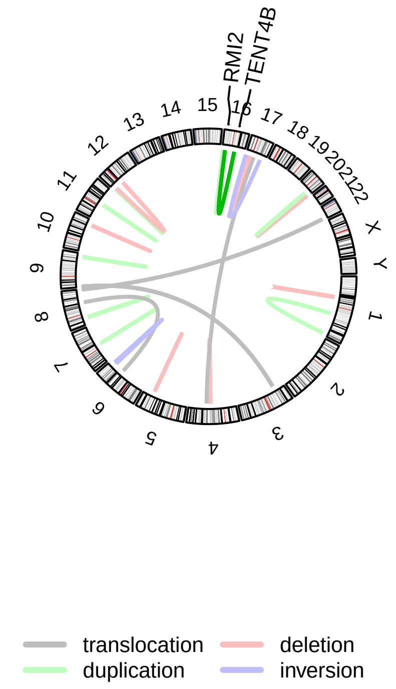


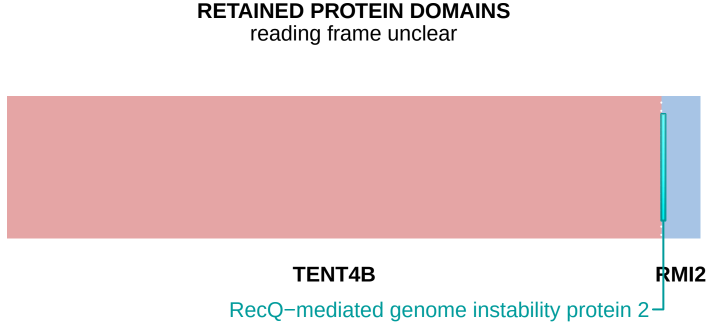


SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0

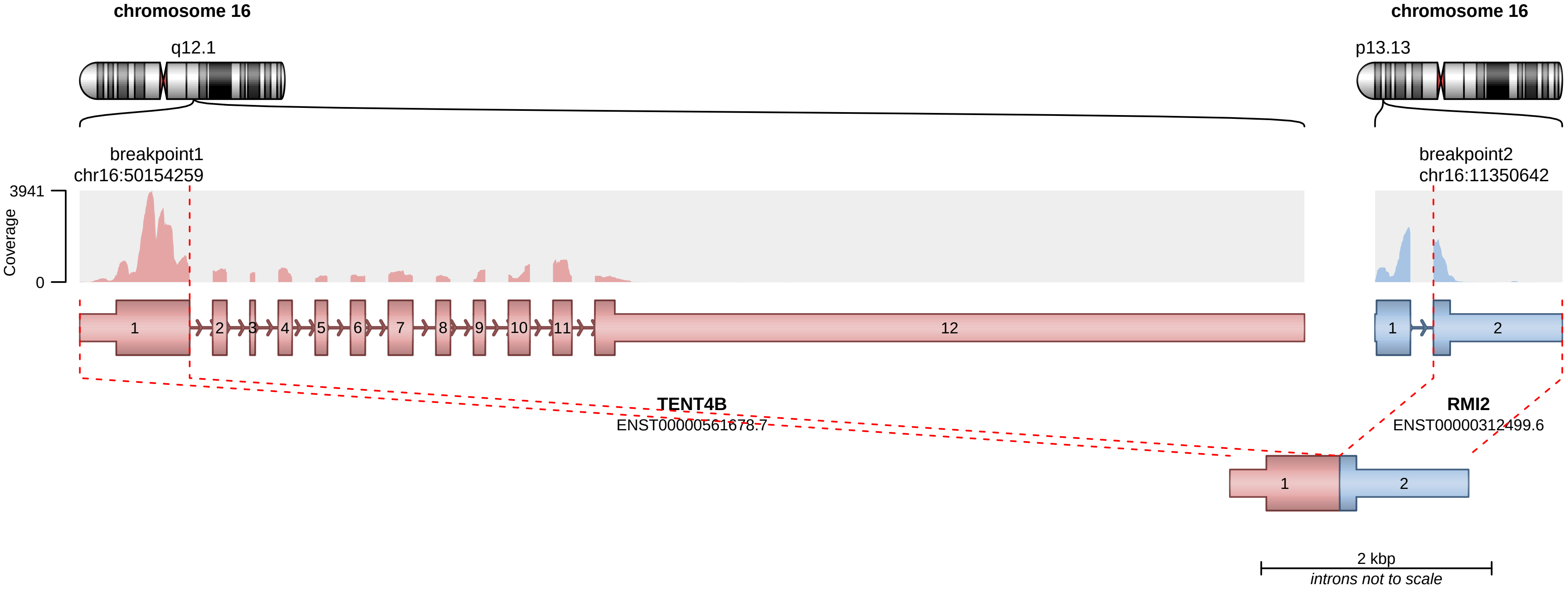


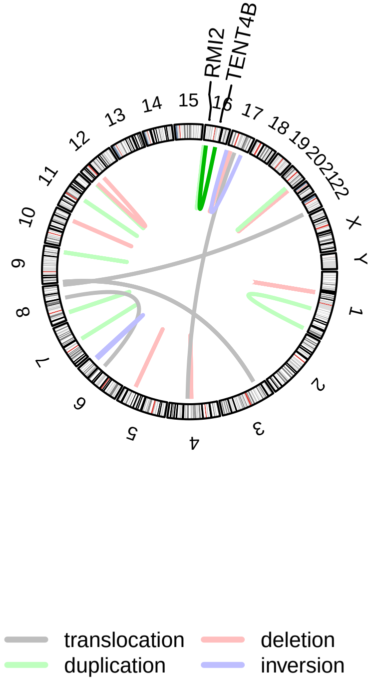


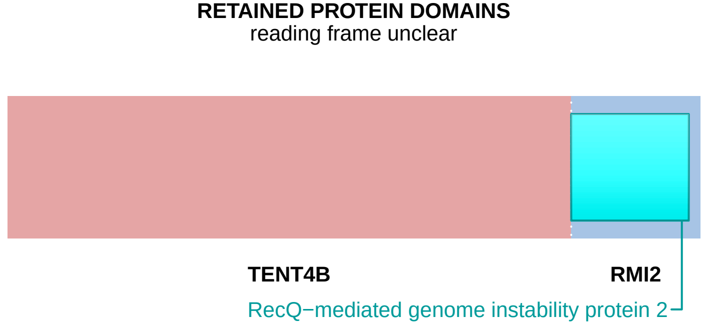


SUPPORTING READ COUNT

Split reads = 46
Discordant mates = 0

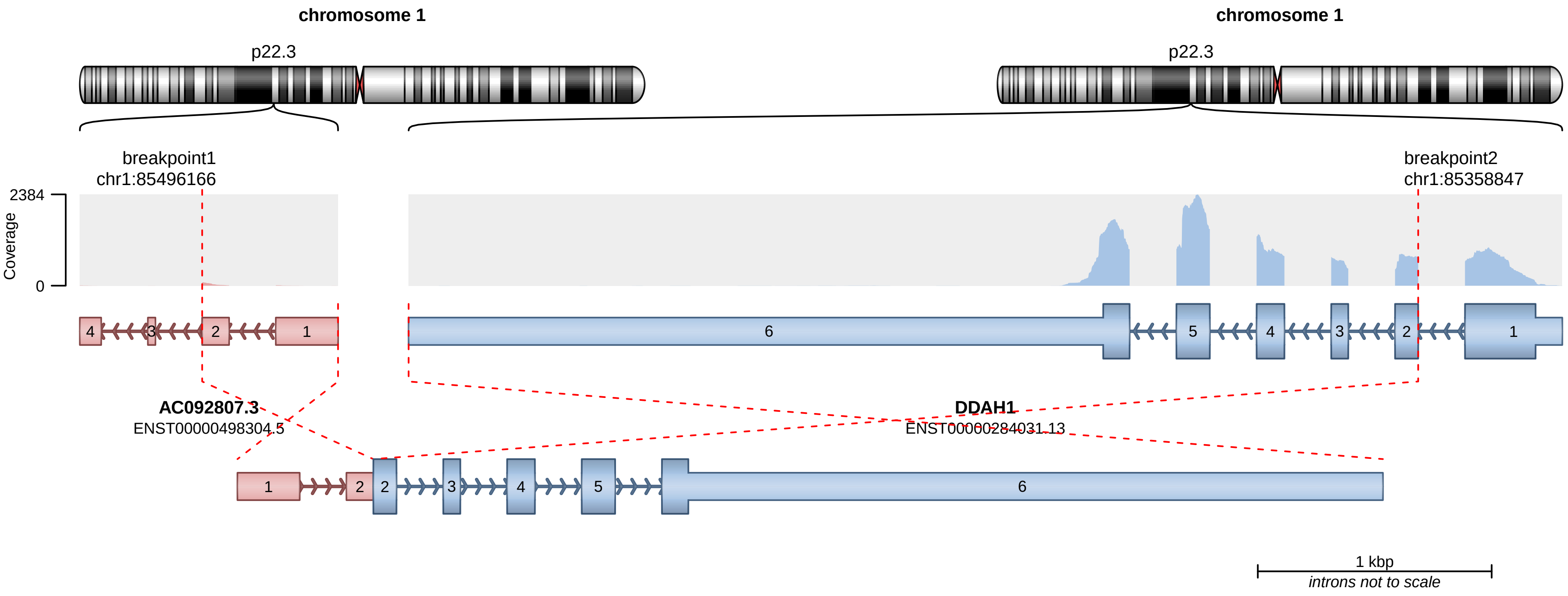


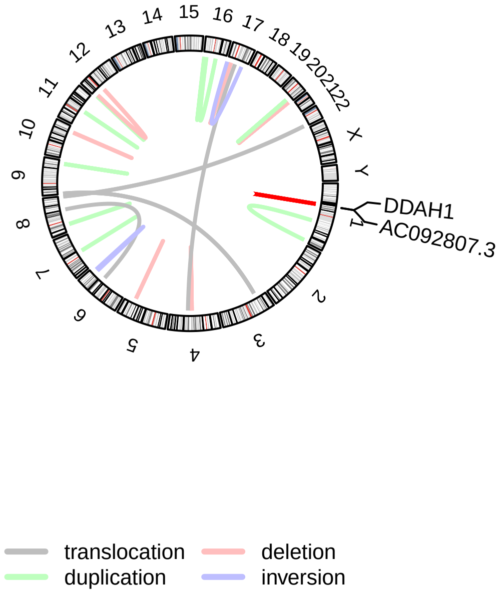


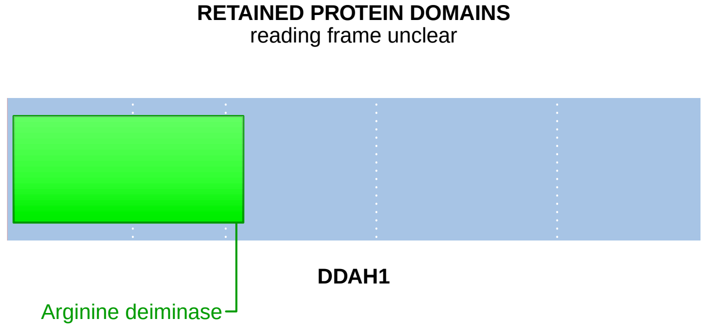


SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

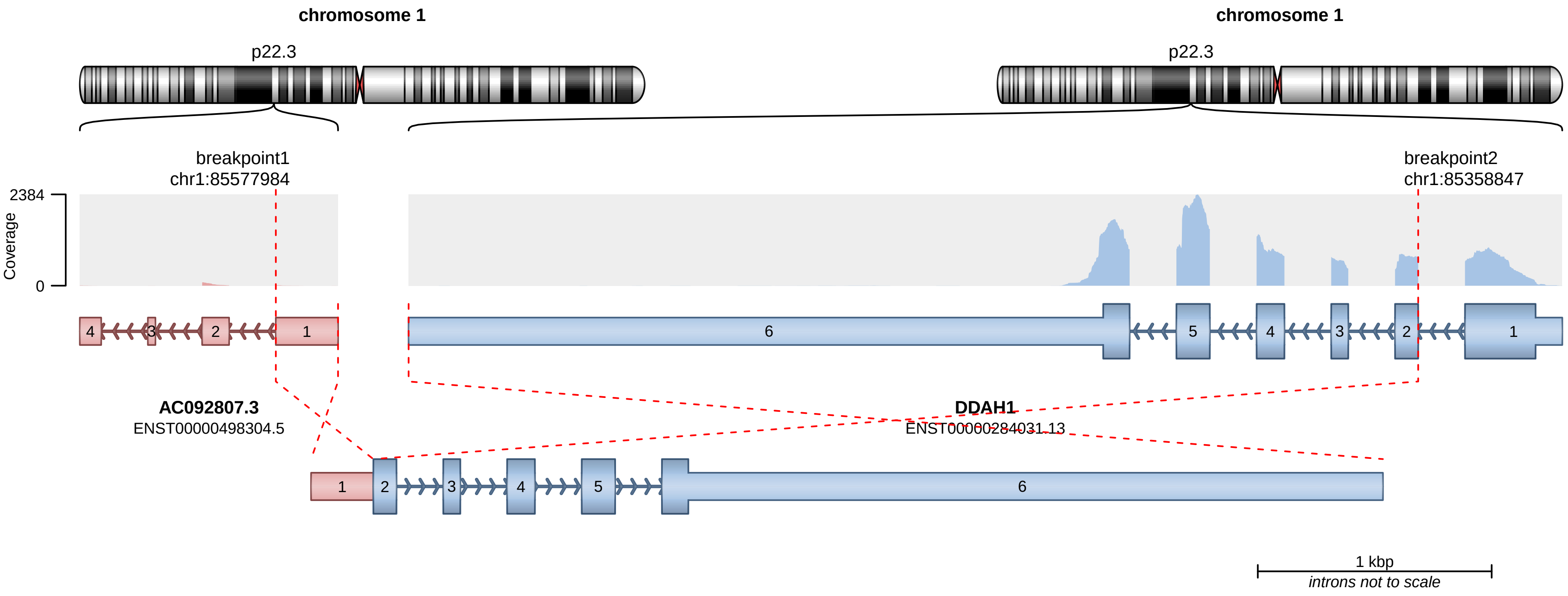


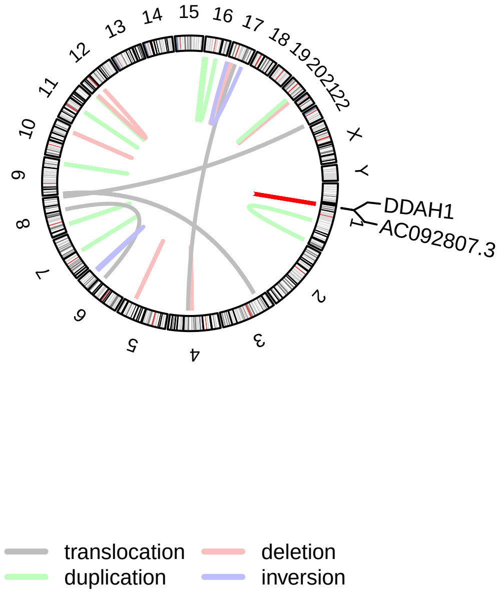




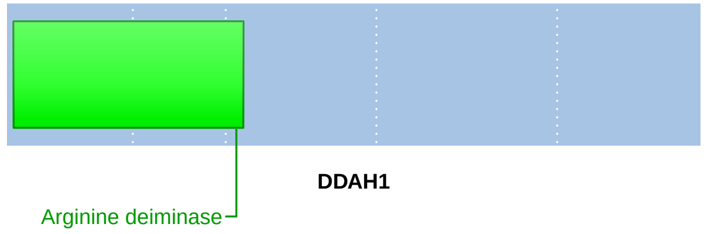
SUPPORTING READ COUNT

Split reads = 41
Discordant mates = 1



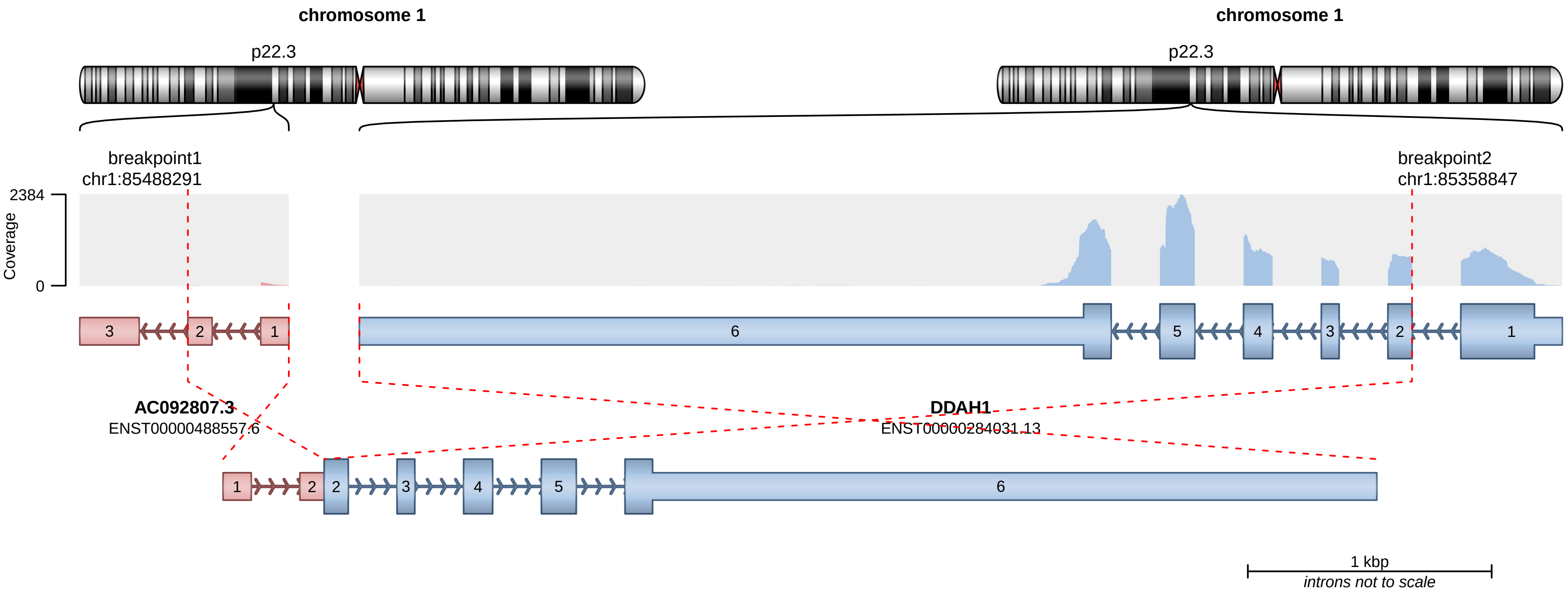


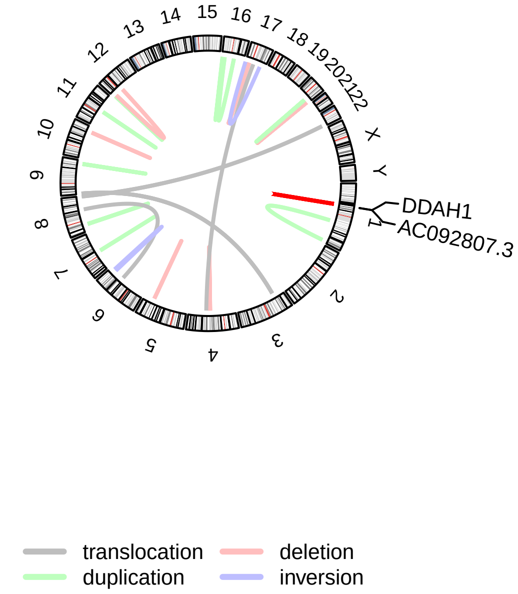
RETAINED PROTEIN DOMAINS
reading frame unclear

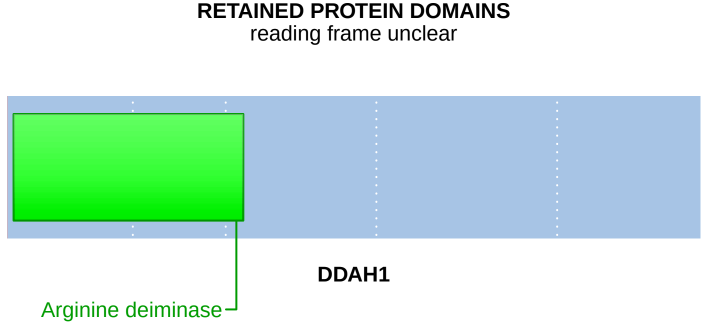


SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

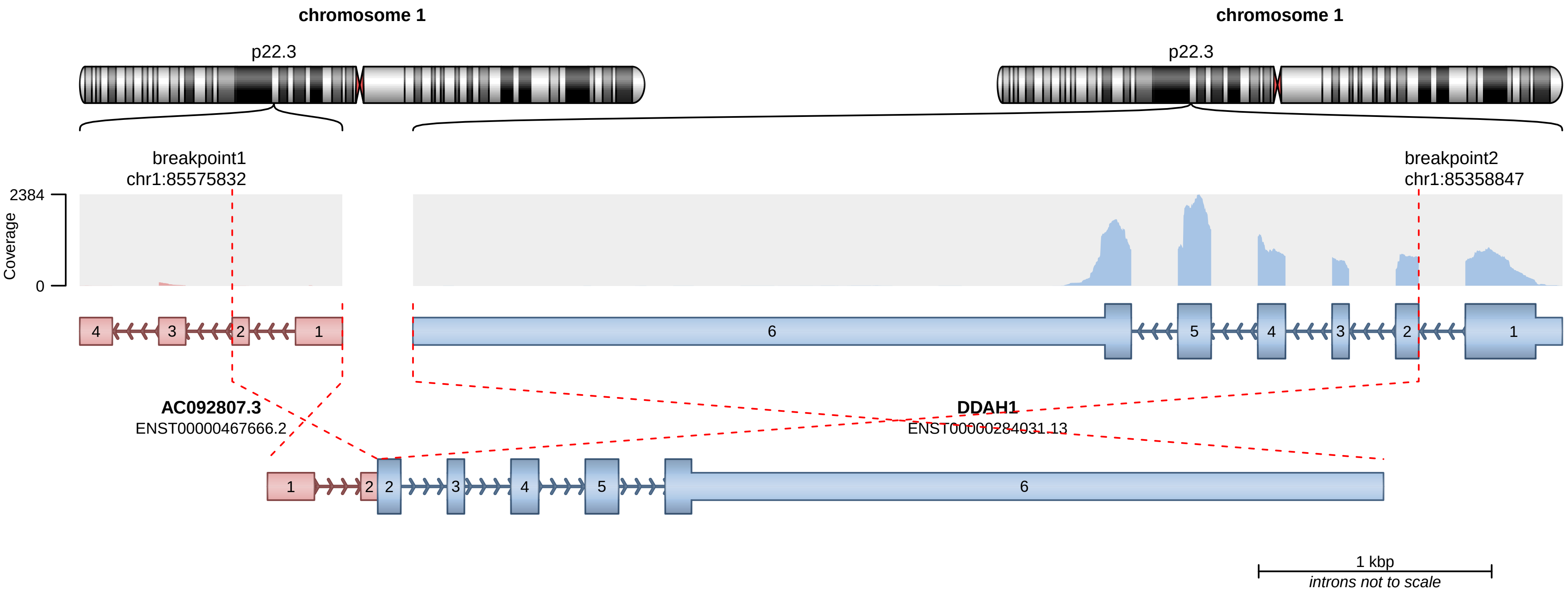


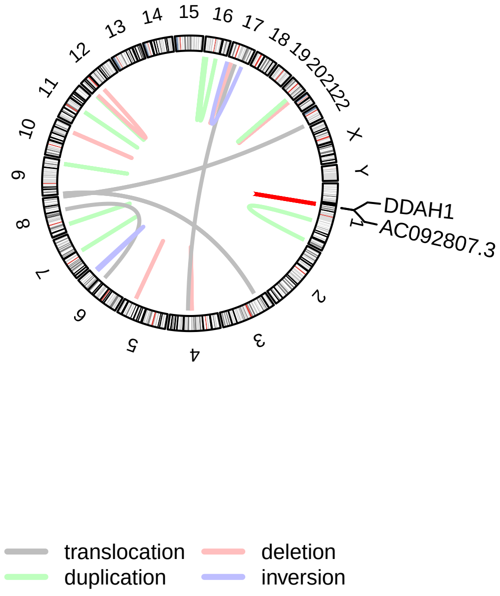


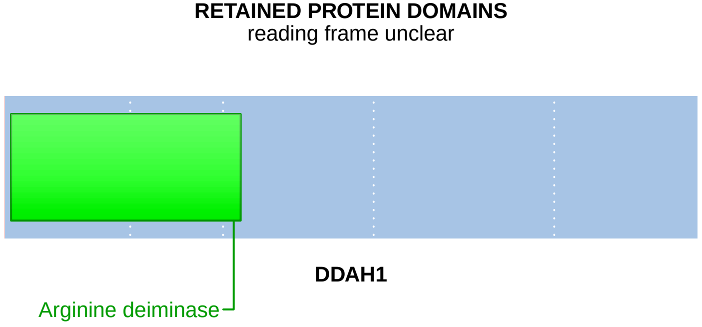


SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 1

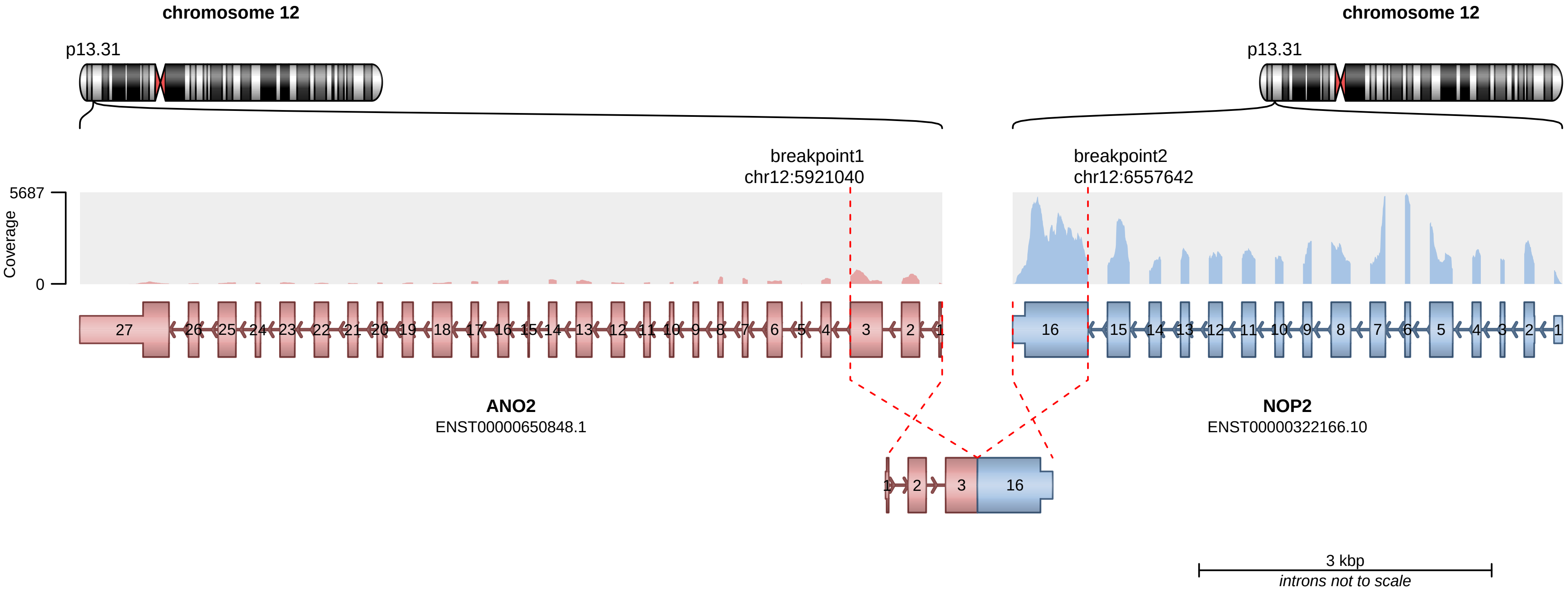


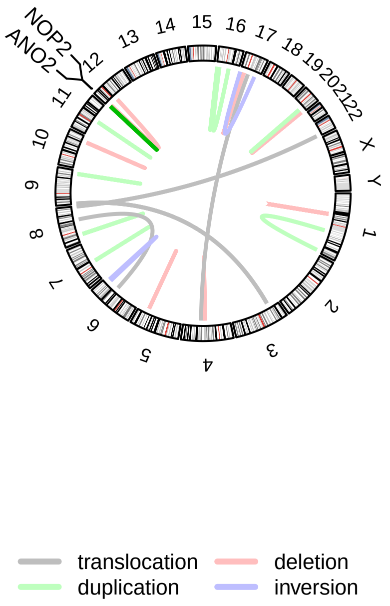


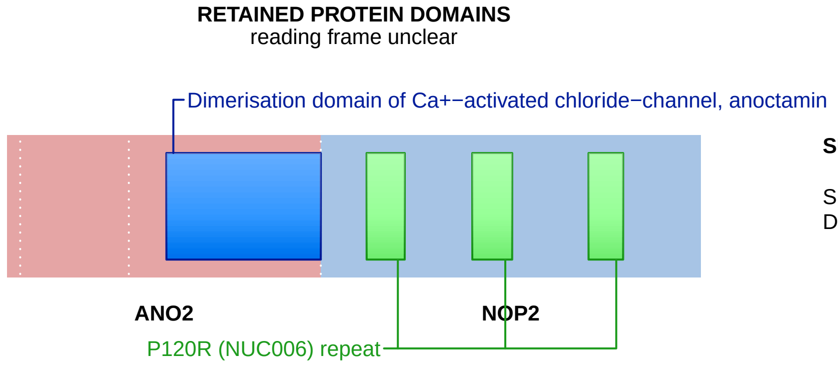


SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

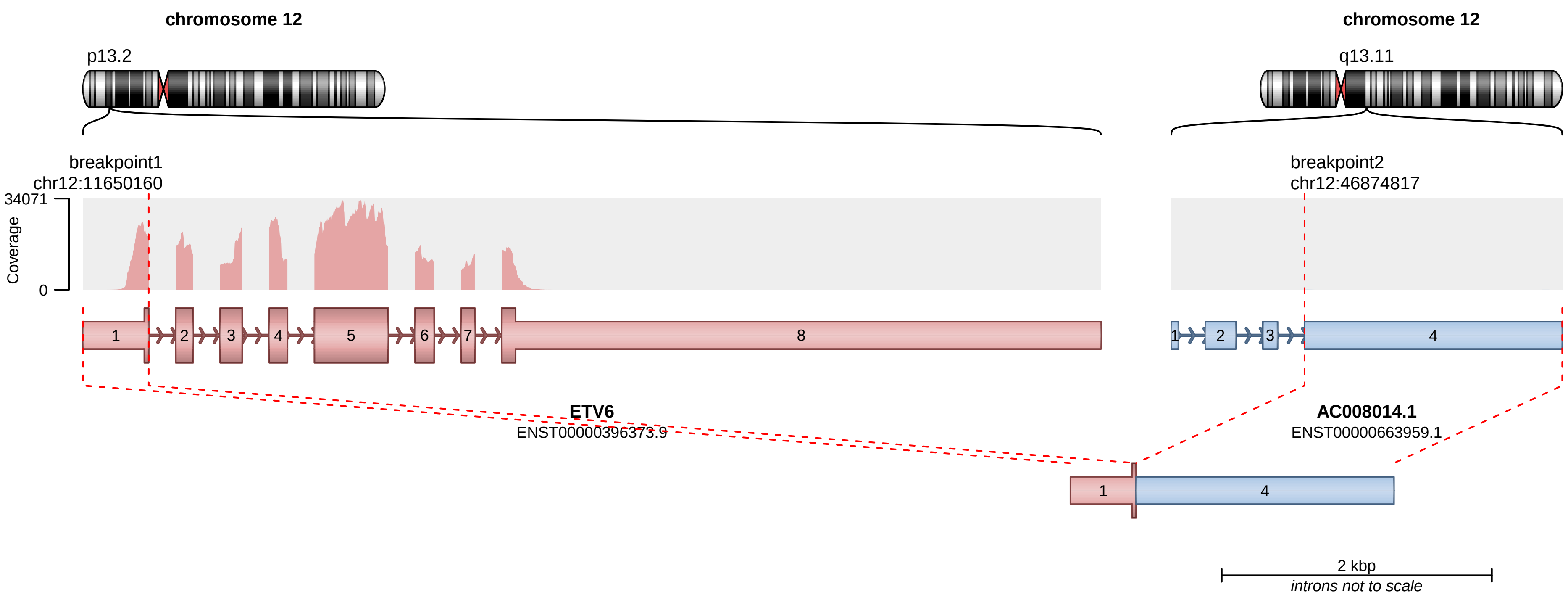


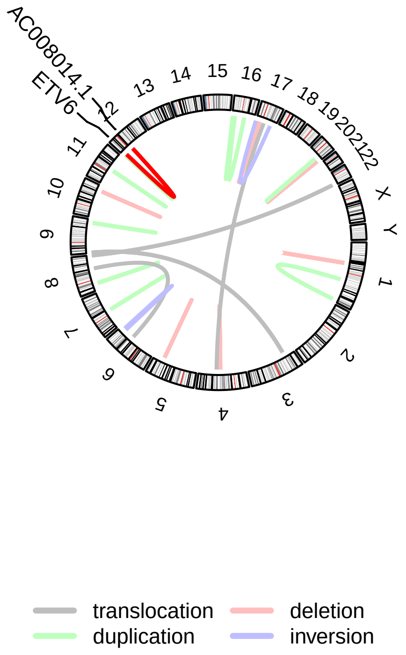




SUPPORTING READ COUNT

Split reads = 33
Discordant mates = 0

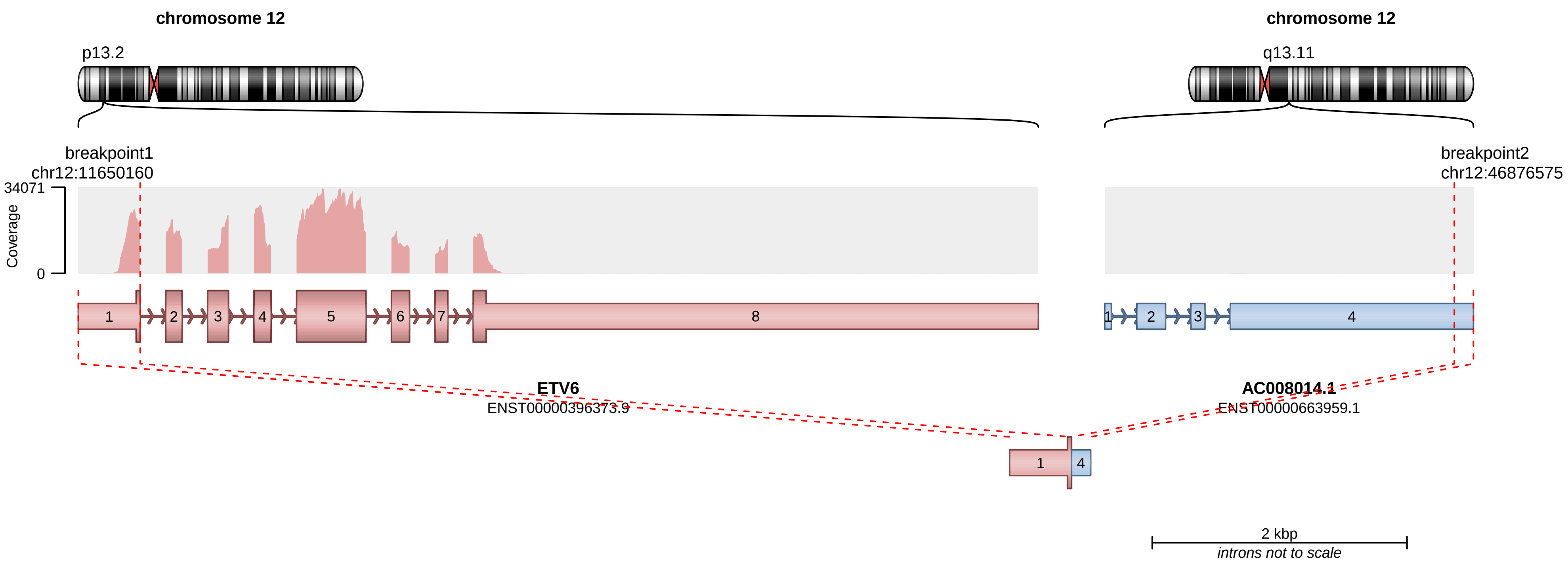


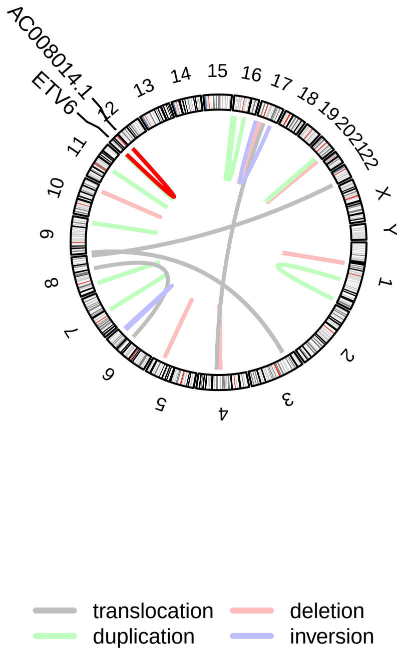


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 31
Discordant mates = 0

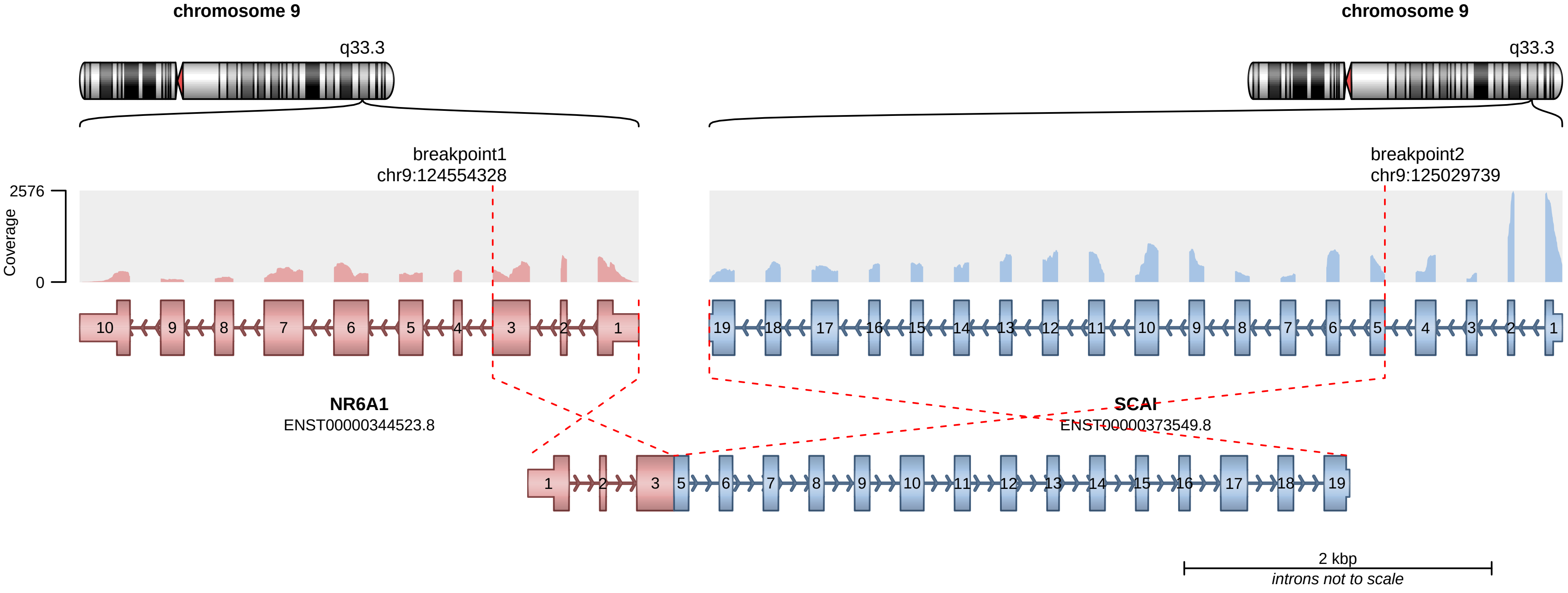


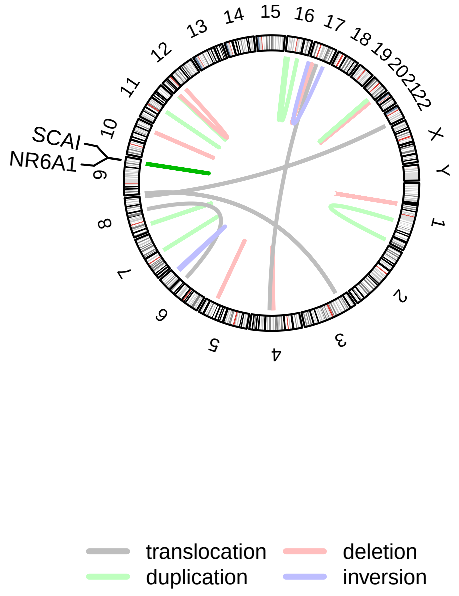


No protein domains retained in fusion.

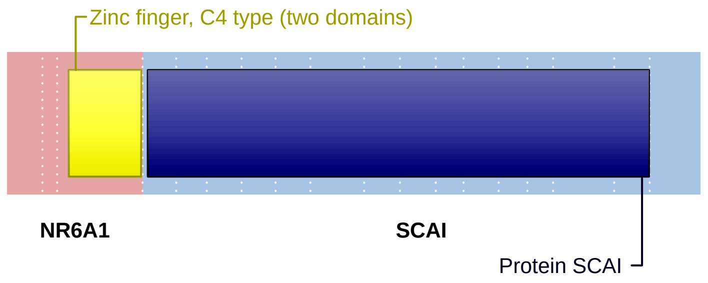
SUPPORTING READ COUNT

Split reads = 27
Discordant mates = 0



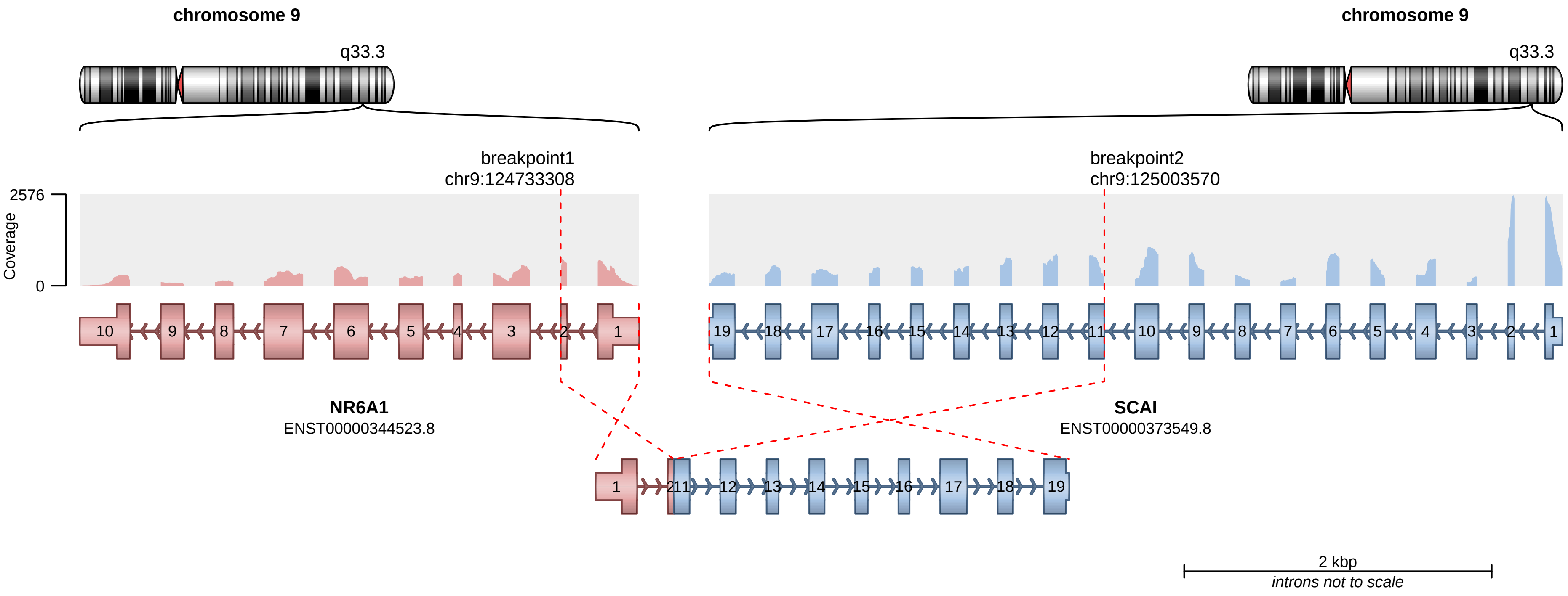


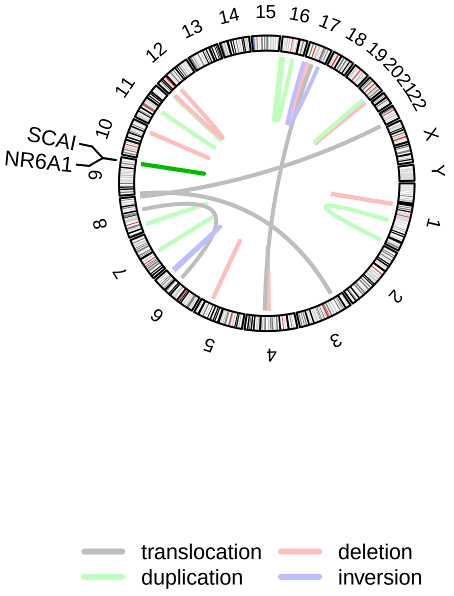
RETAINED PROTEIN DOMAINS
reading frame unclear

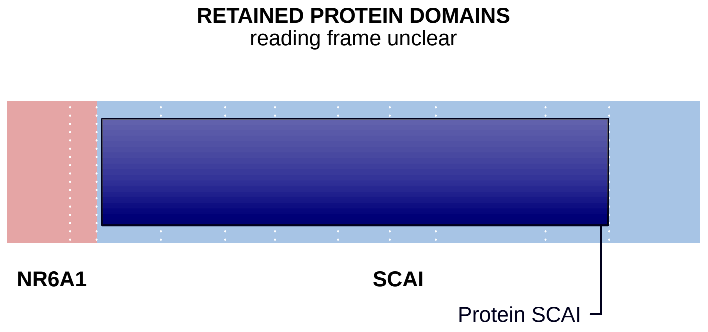


SUPPORTING READ COUNT

Split reads = 23
Discordant mates = 0

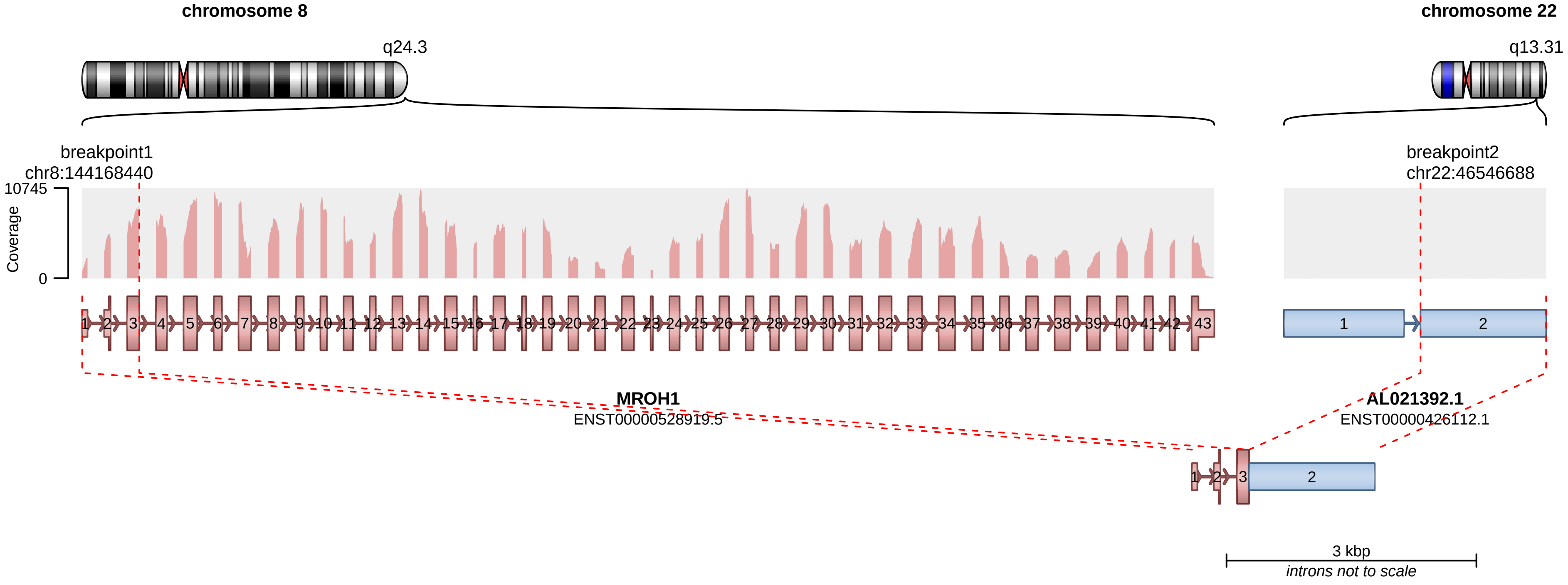


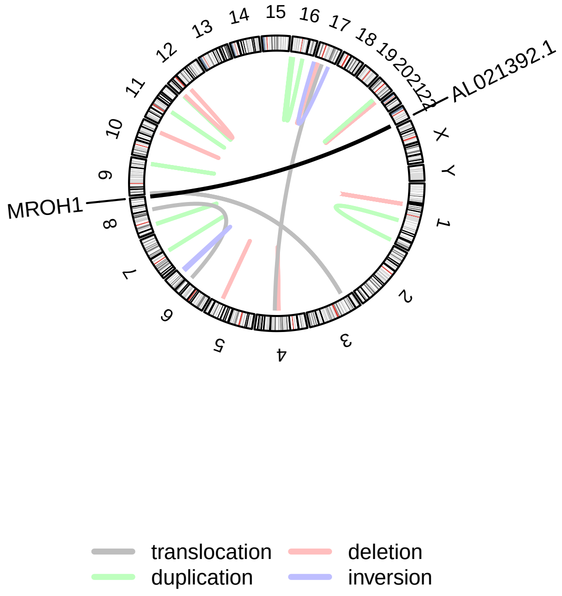




SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

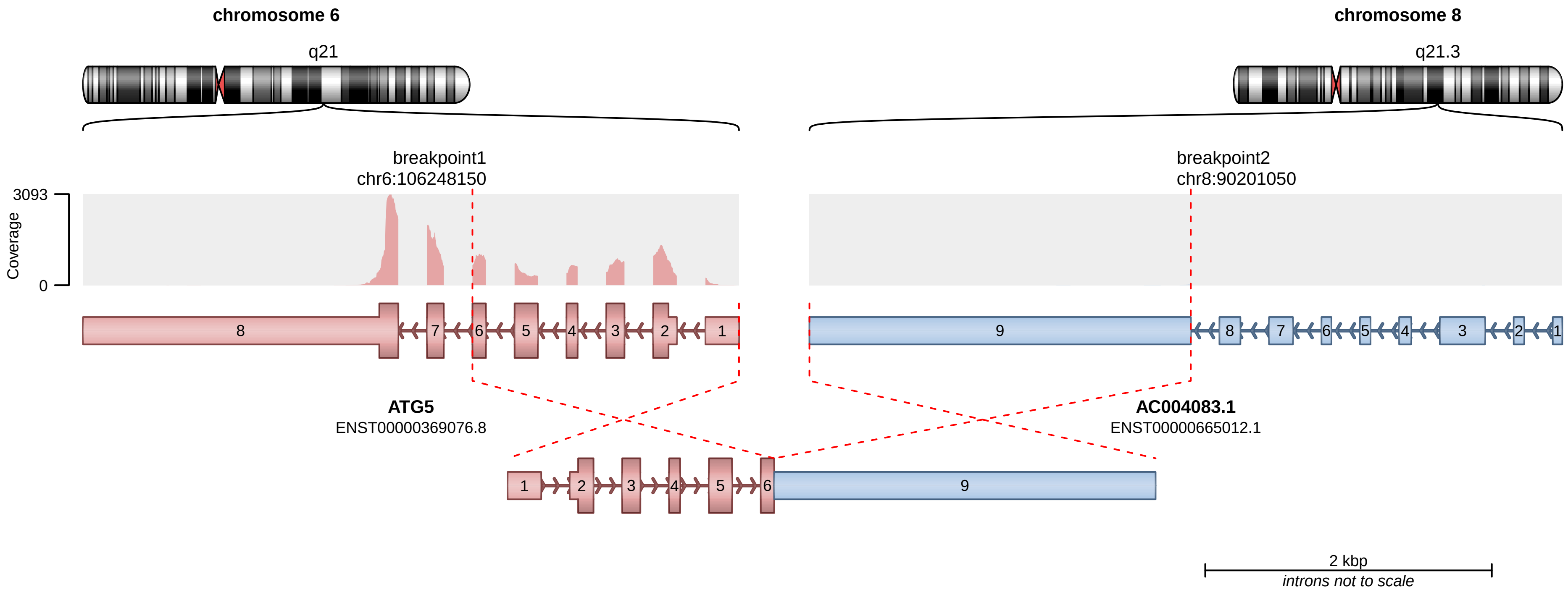


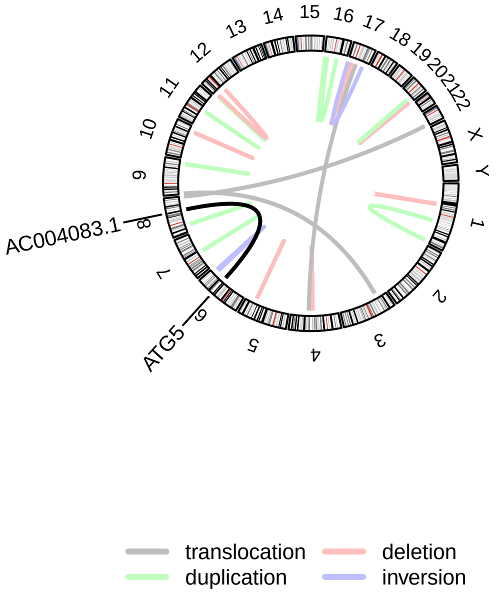


No protein domains retained in fusion.

SUPPORTING READ COUNT

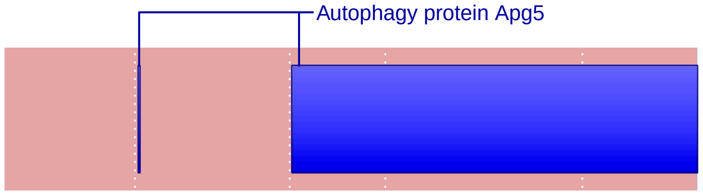
Split reads = 21
Discordant mates = 0





RETAINED PROTEIN DOMAINS

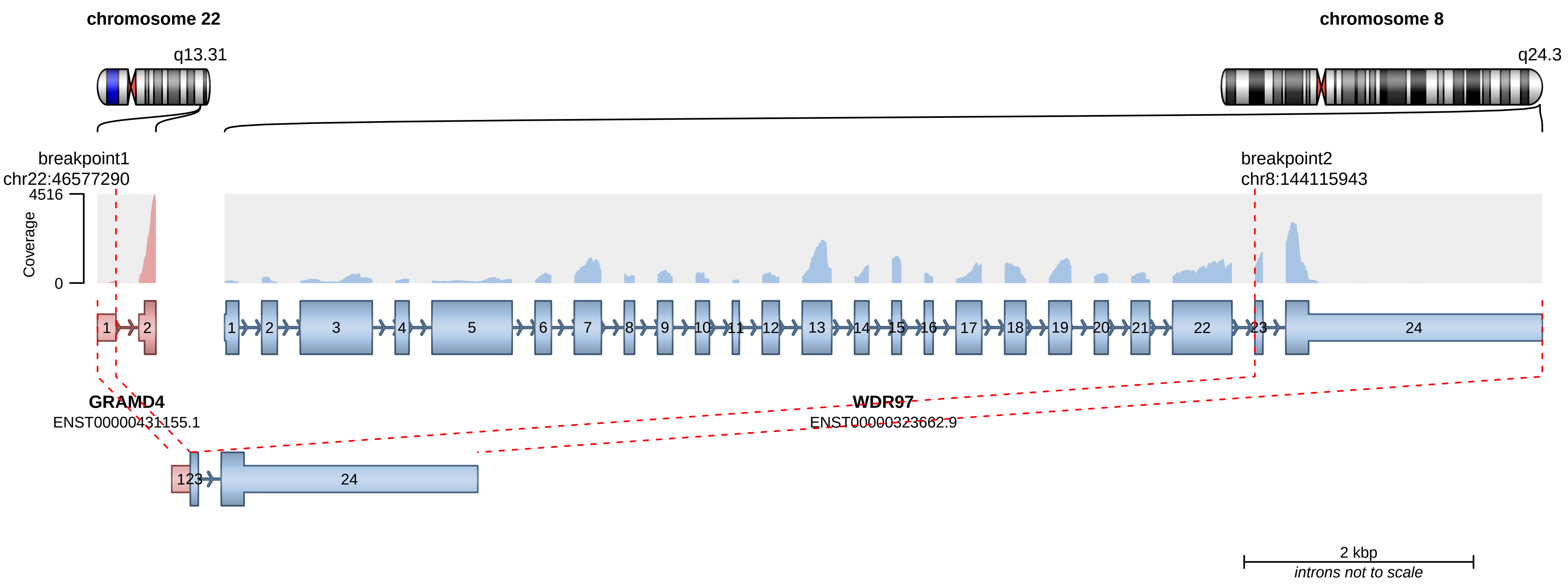
reading frame unclear

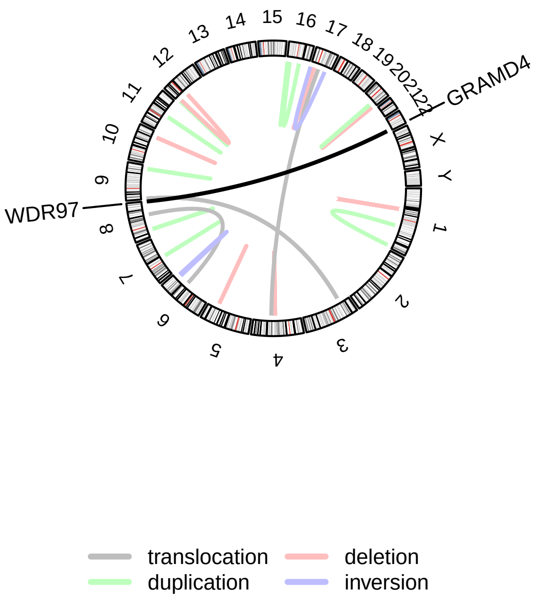


ATG5

SUPPORTING READ COUNT

Split reads = 19
Discordant mates = 0

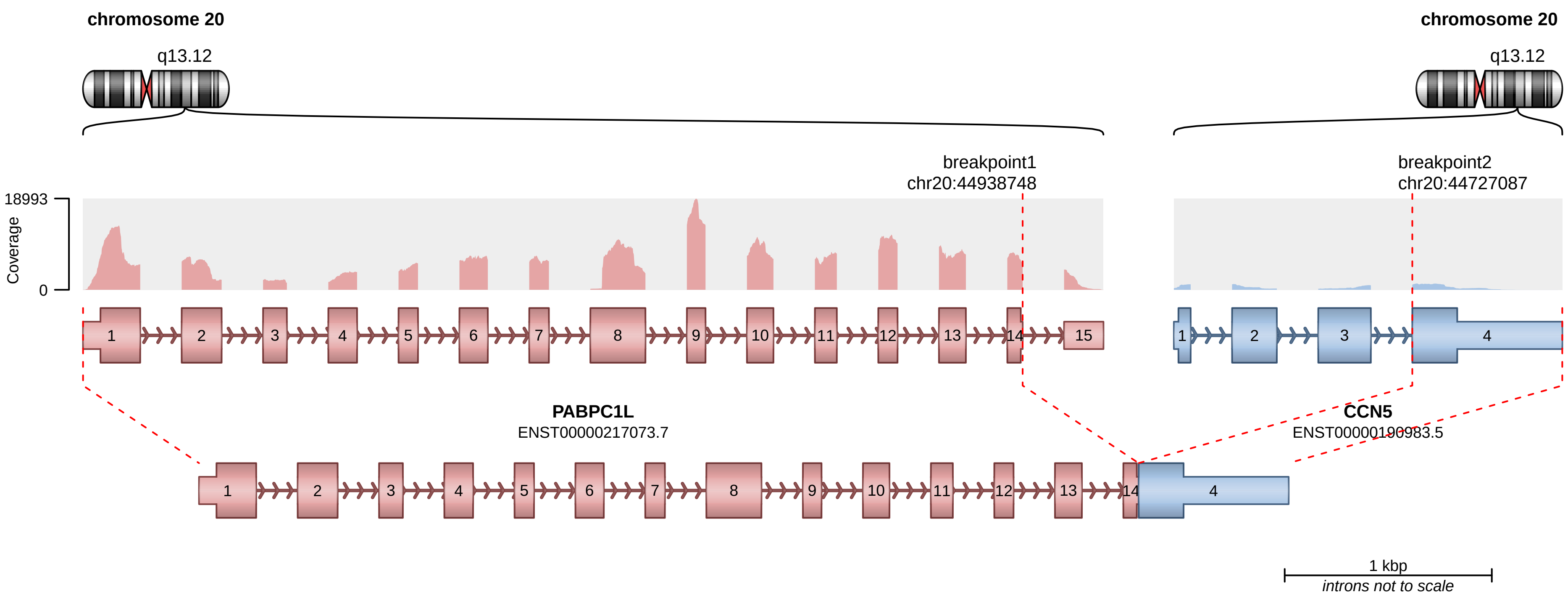


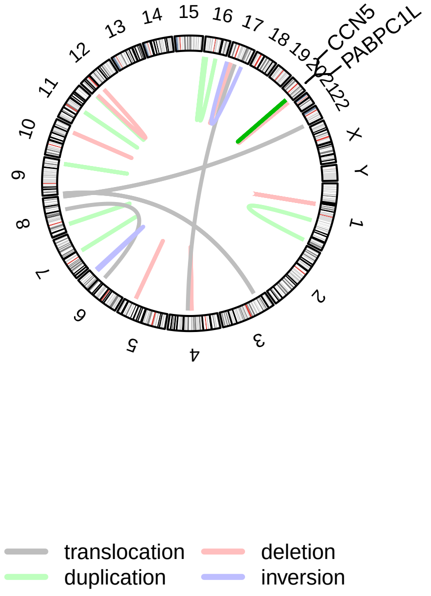


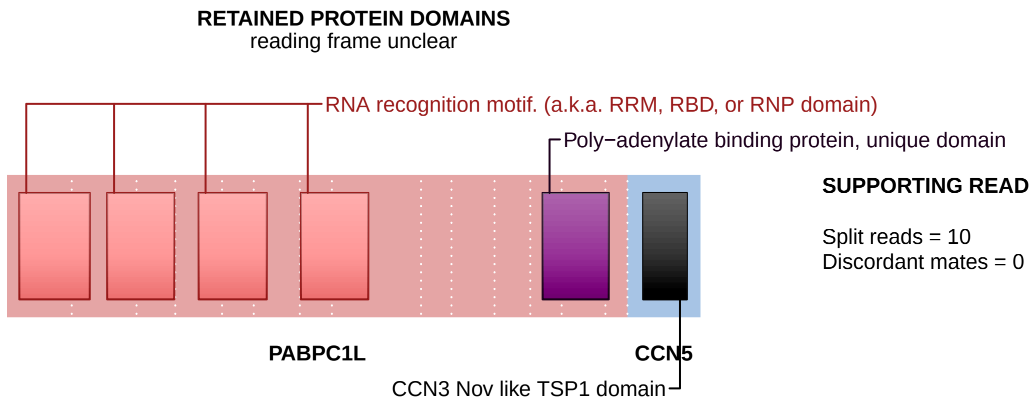
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 19
Discordant mates = 0

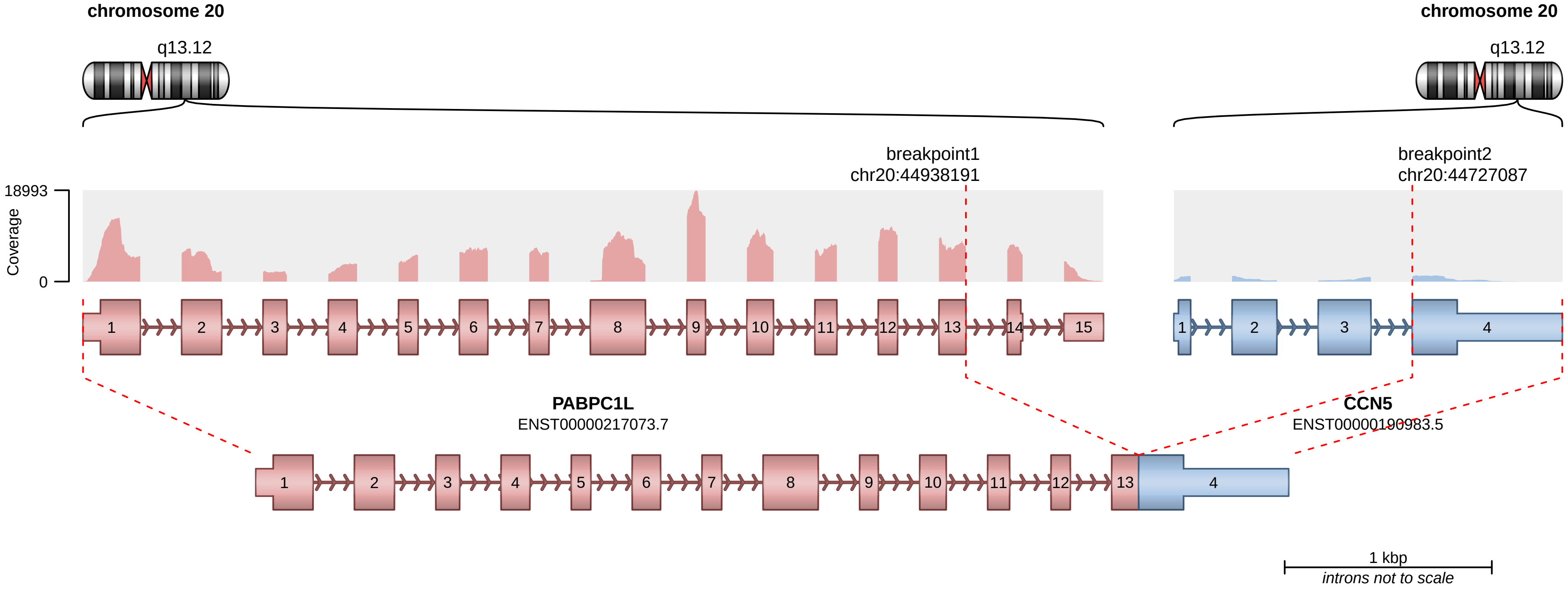


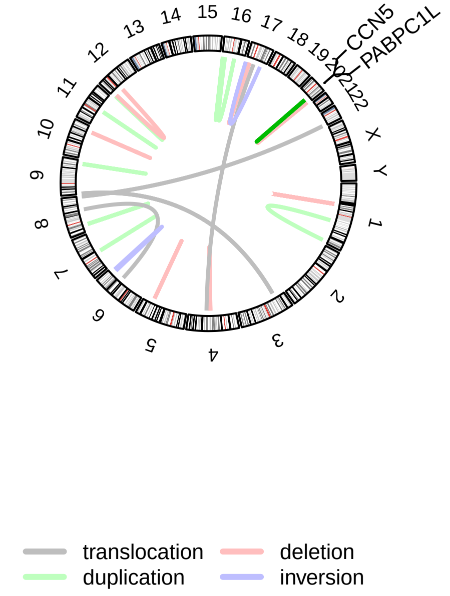


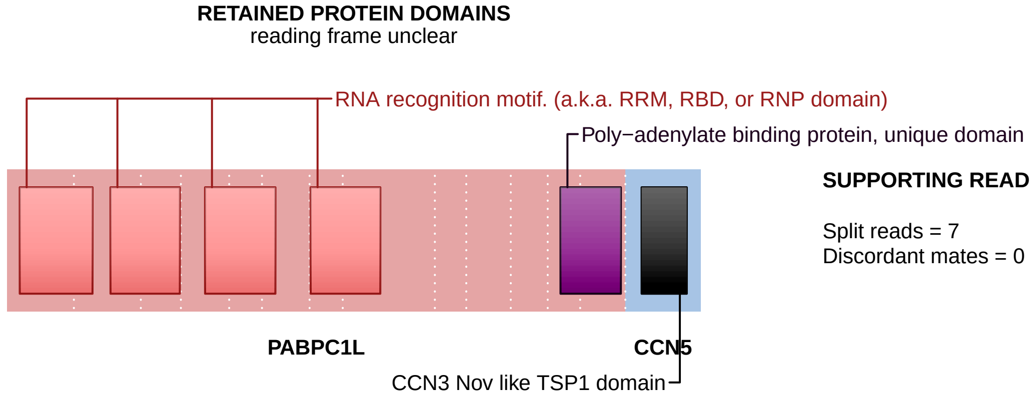


SUPPORTING READ COUNT

Split reads = 10
Discordant mates = 0

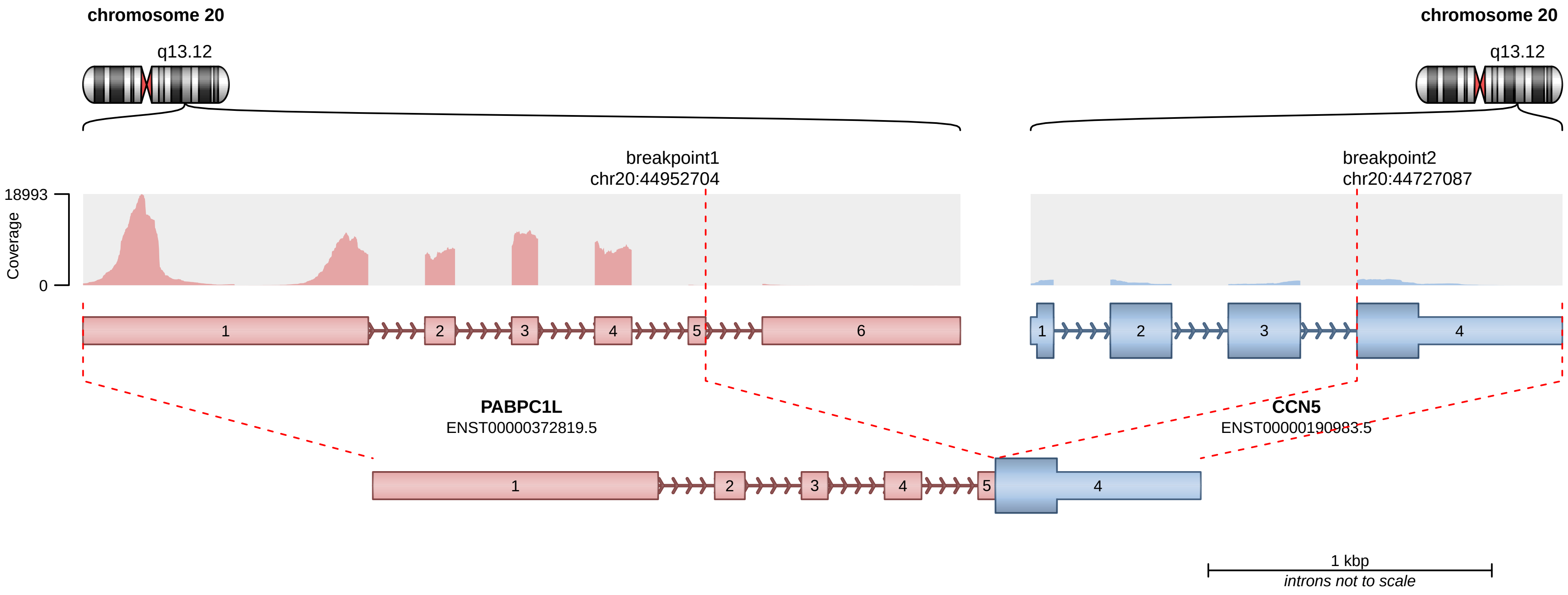


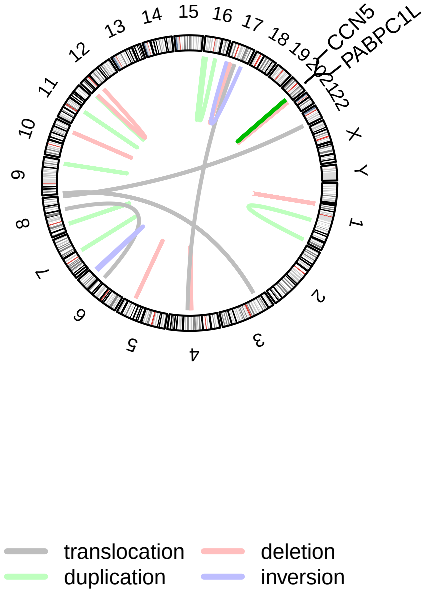




SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

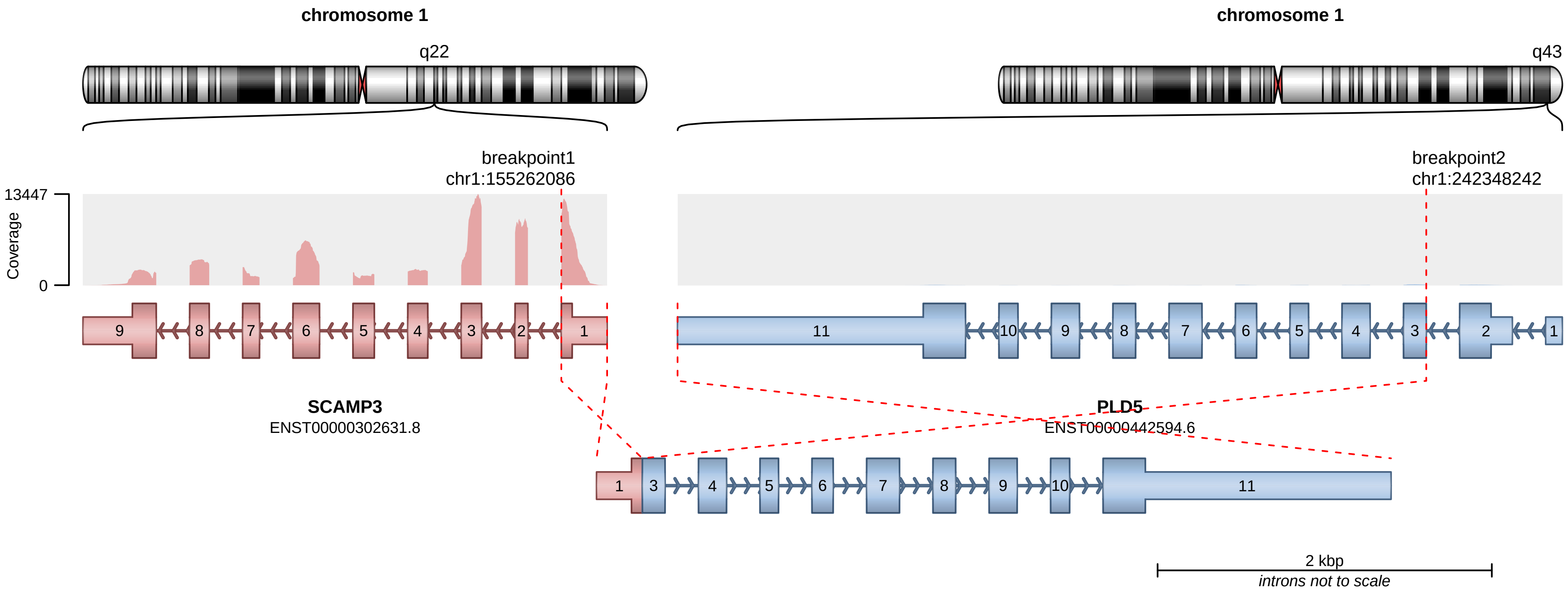


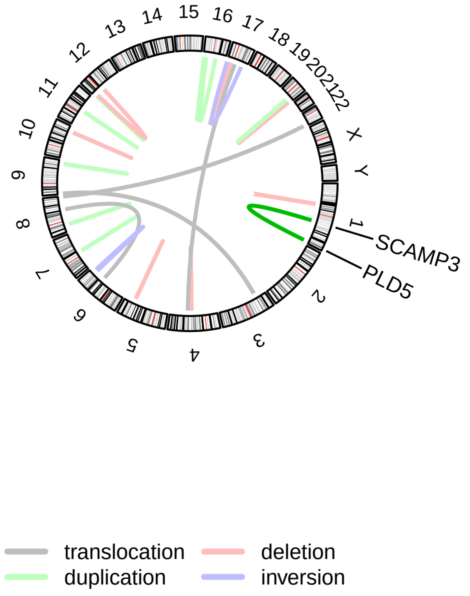


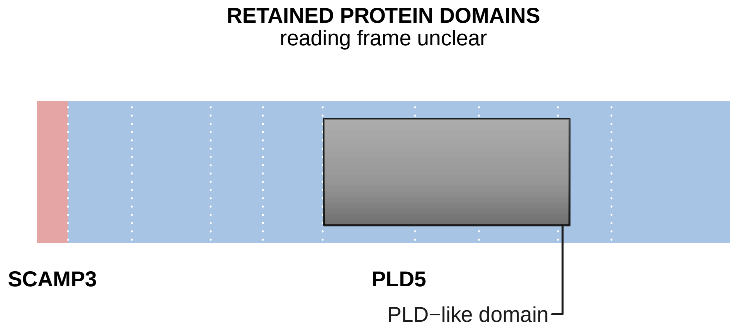
RETAINED PROTEIN DOMAINS
reading frame unclear

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

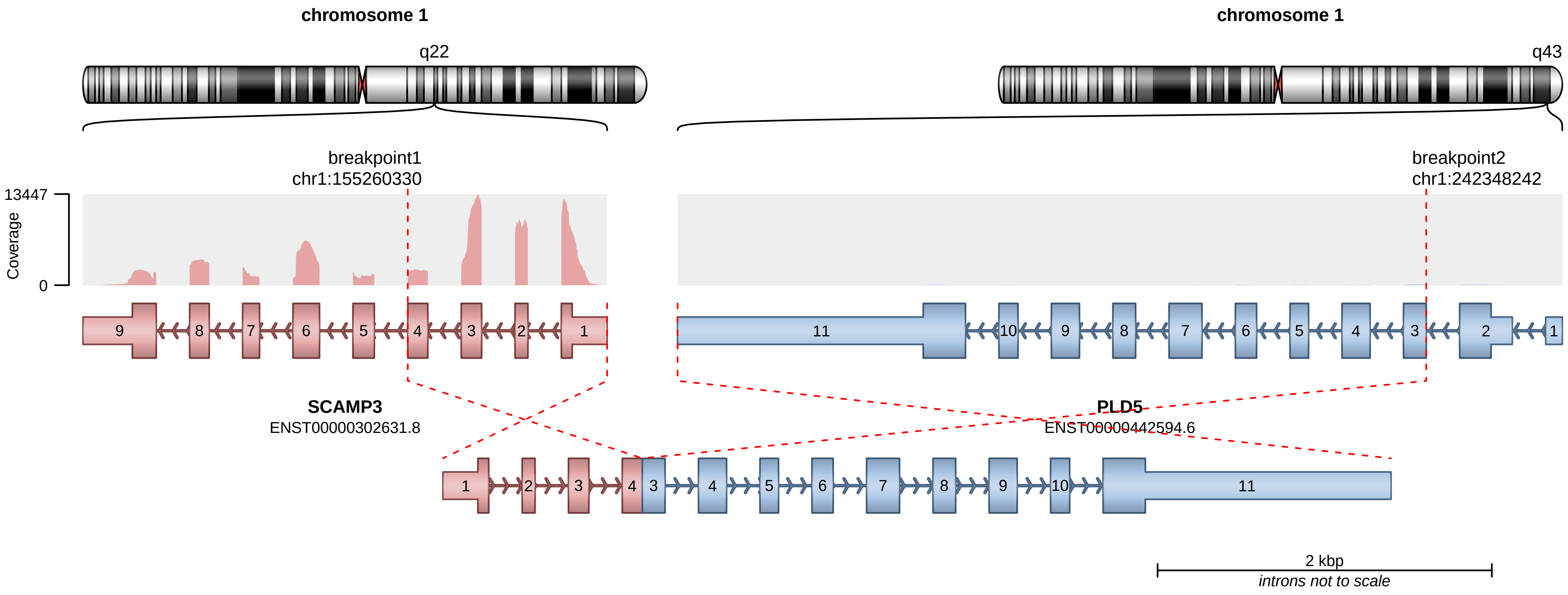


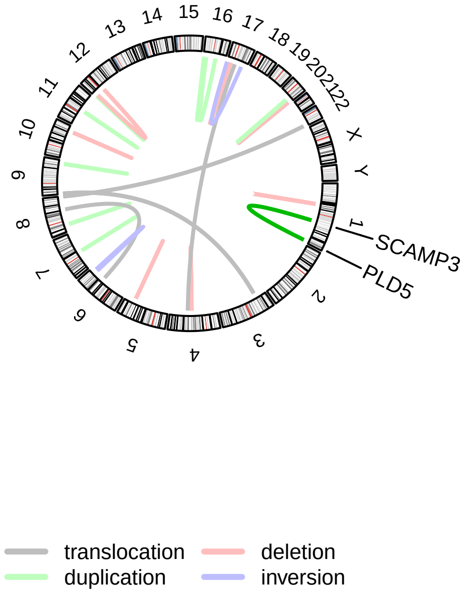


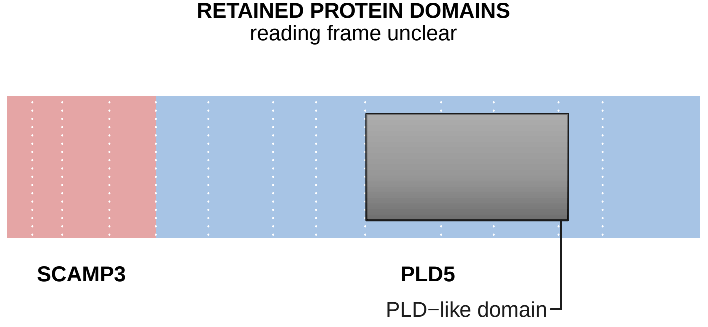


SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 1

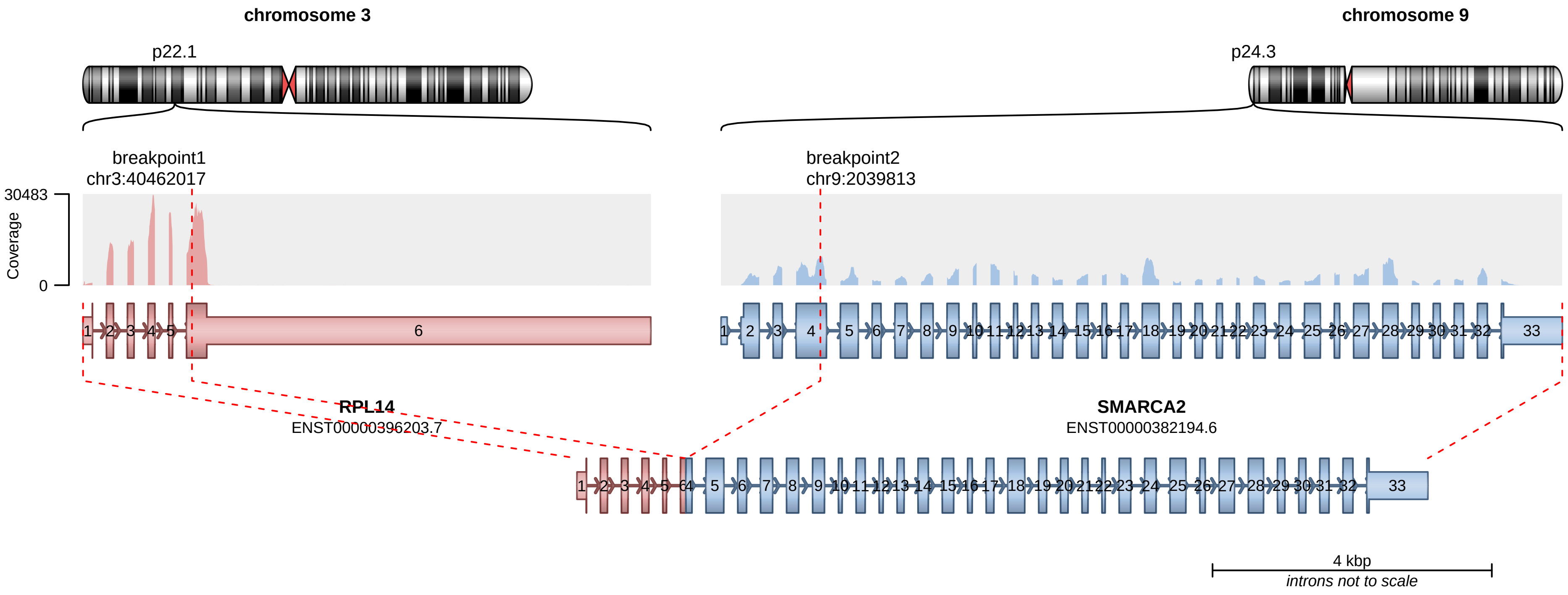


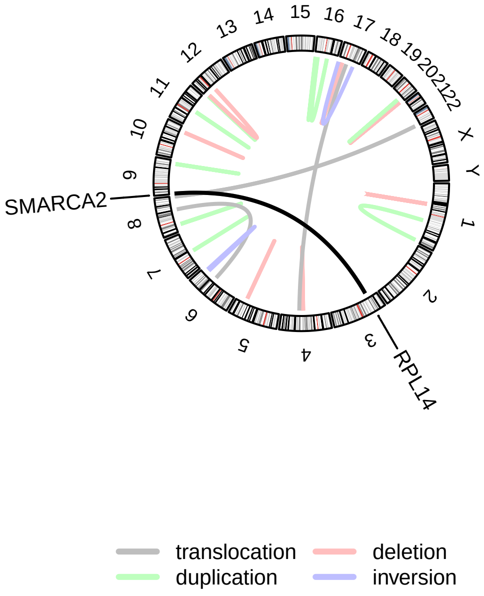


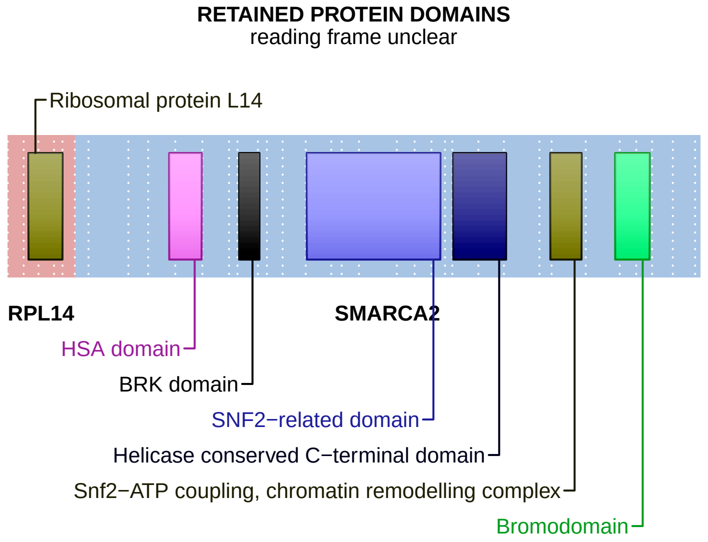


SUPPORTING READ COUNT

Split reads = 5
Discordant mates = 2

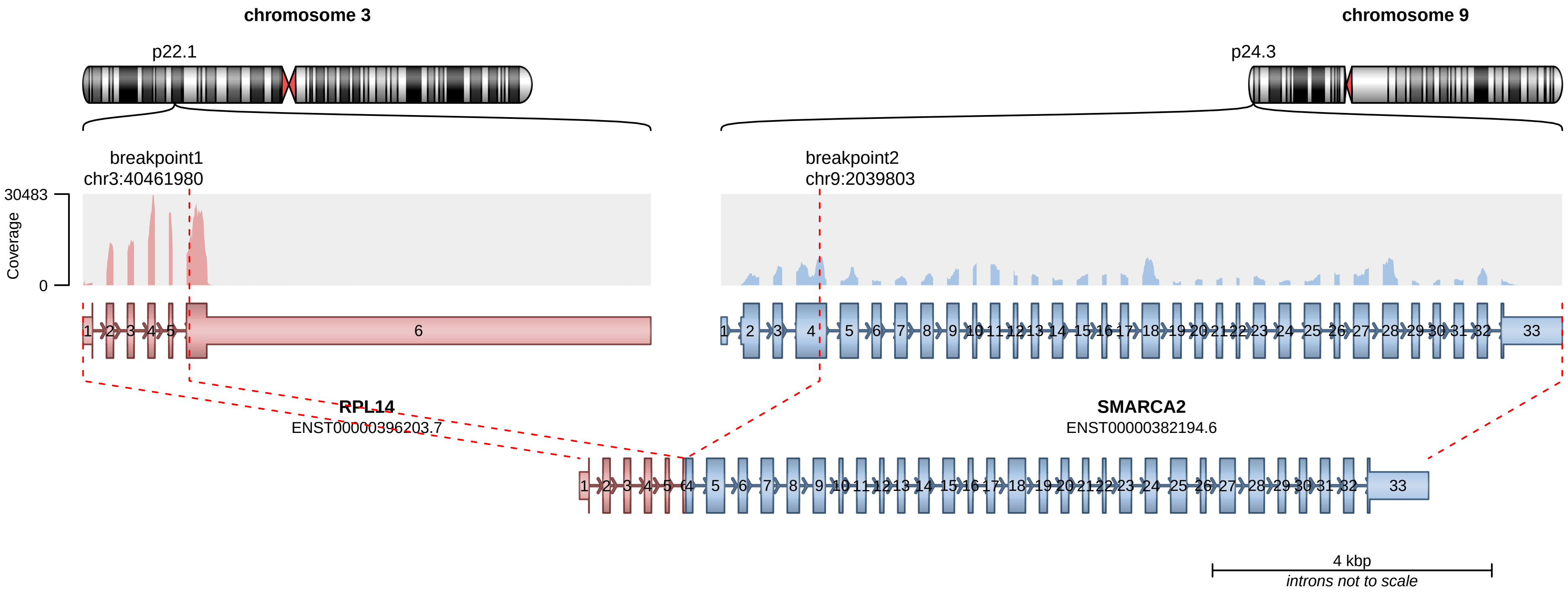


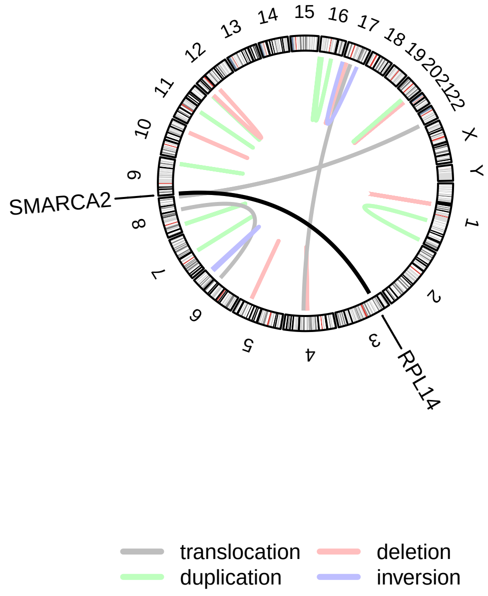


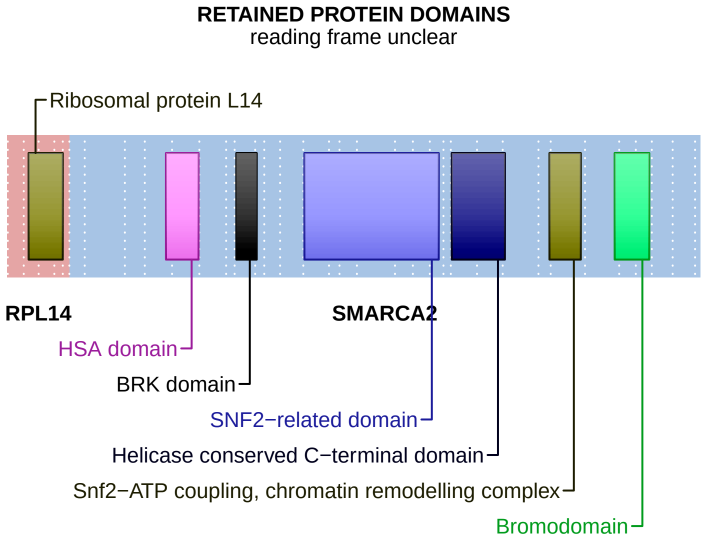


SUPPORTING READ COUNT

Split reads = 4
Discordant mates = 0

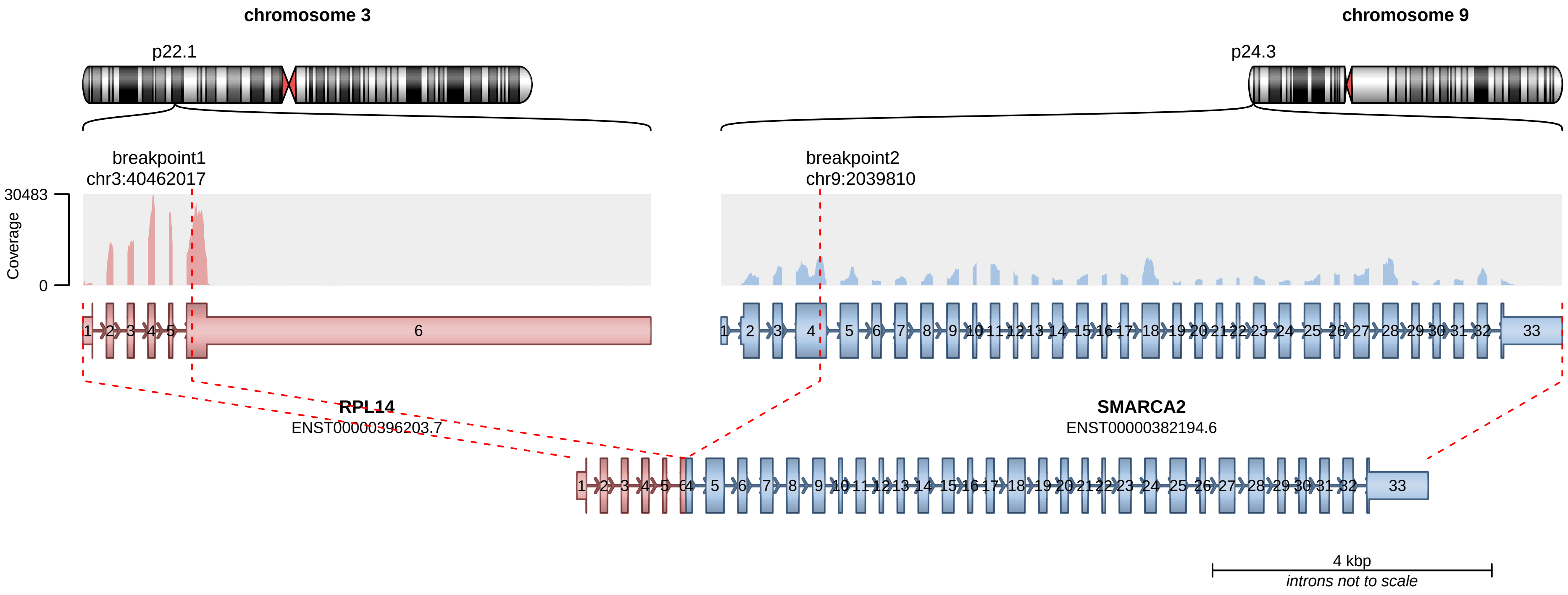


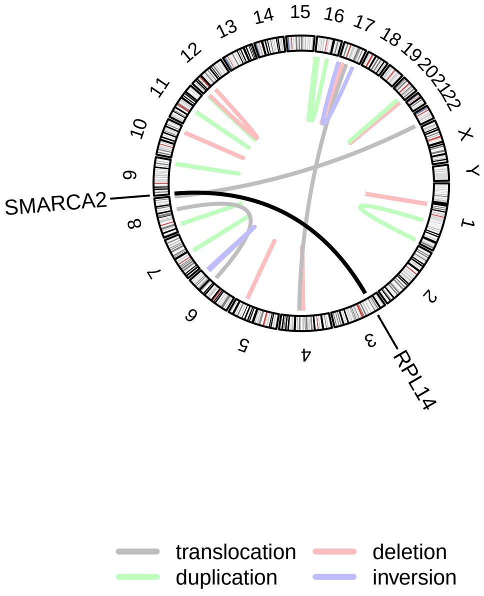


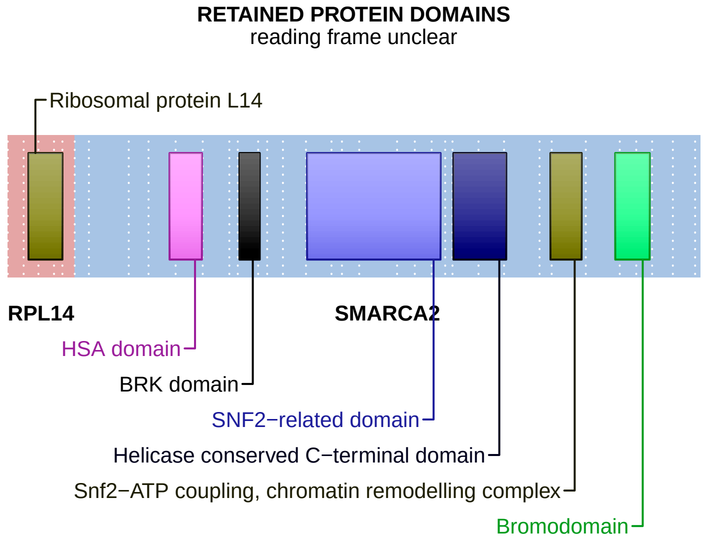


SUPPORTING READ COUNT

Split reads = 4
Discordant mates = 0

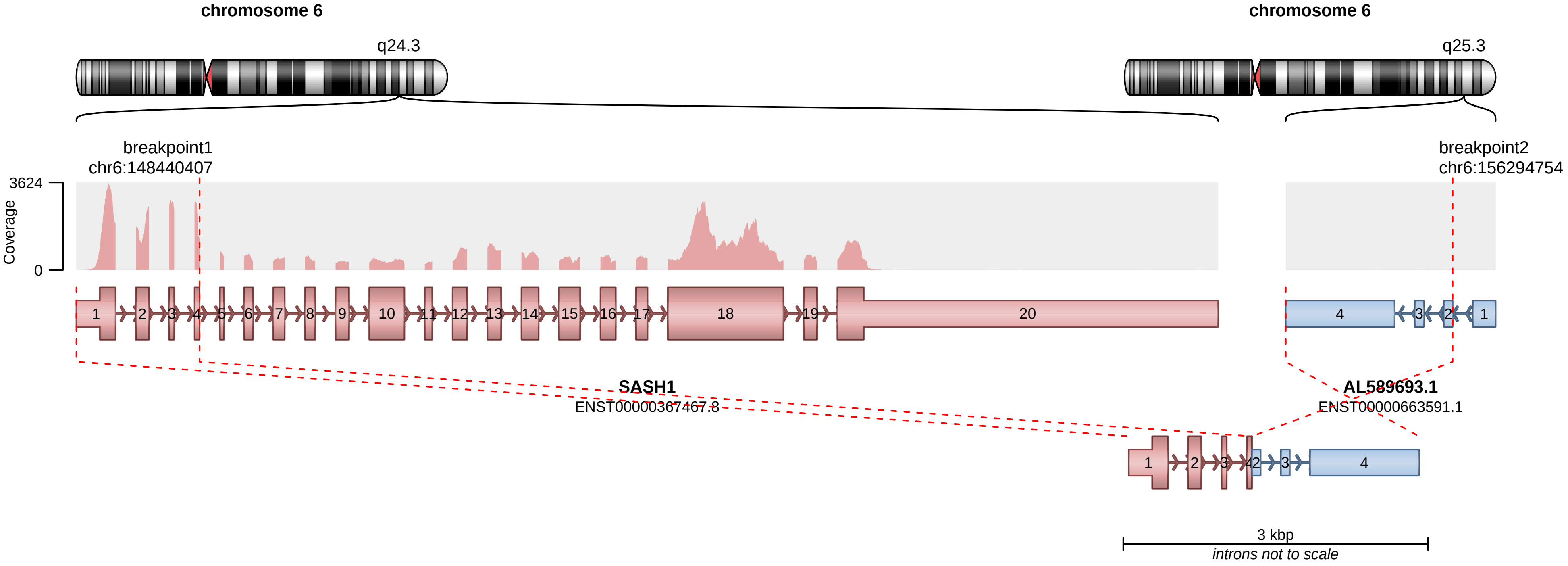


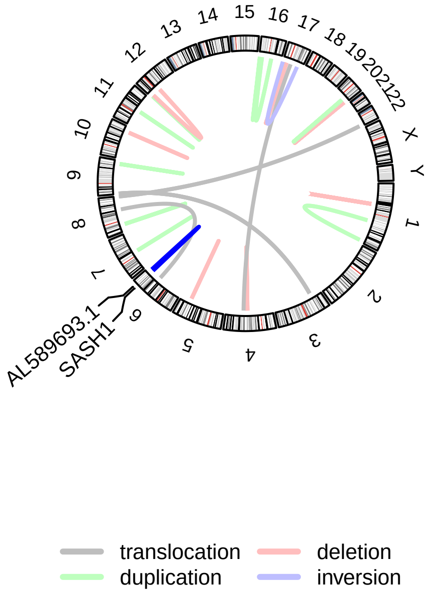




SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0





No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0