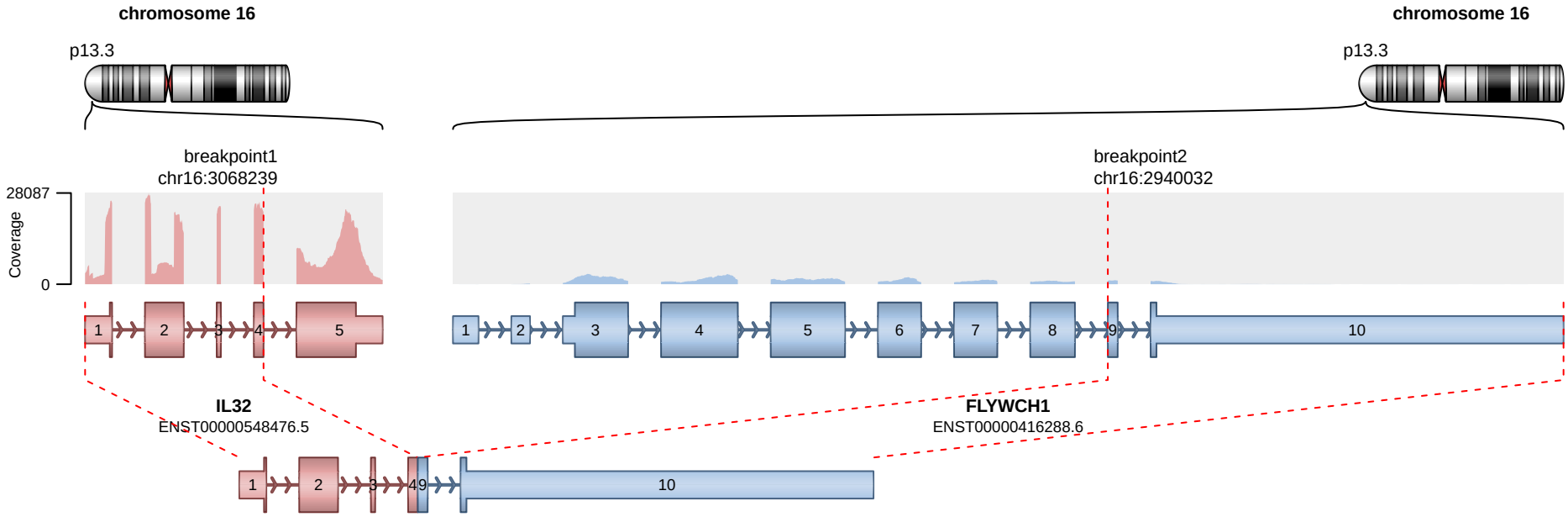
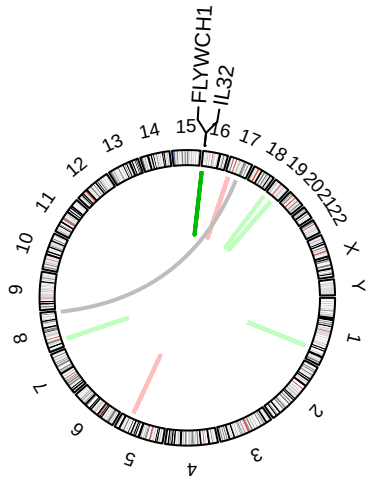
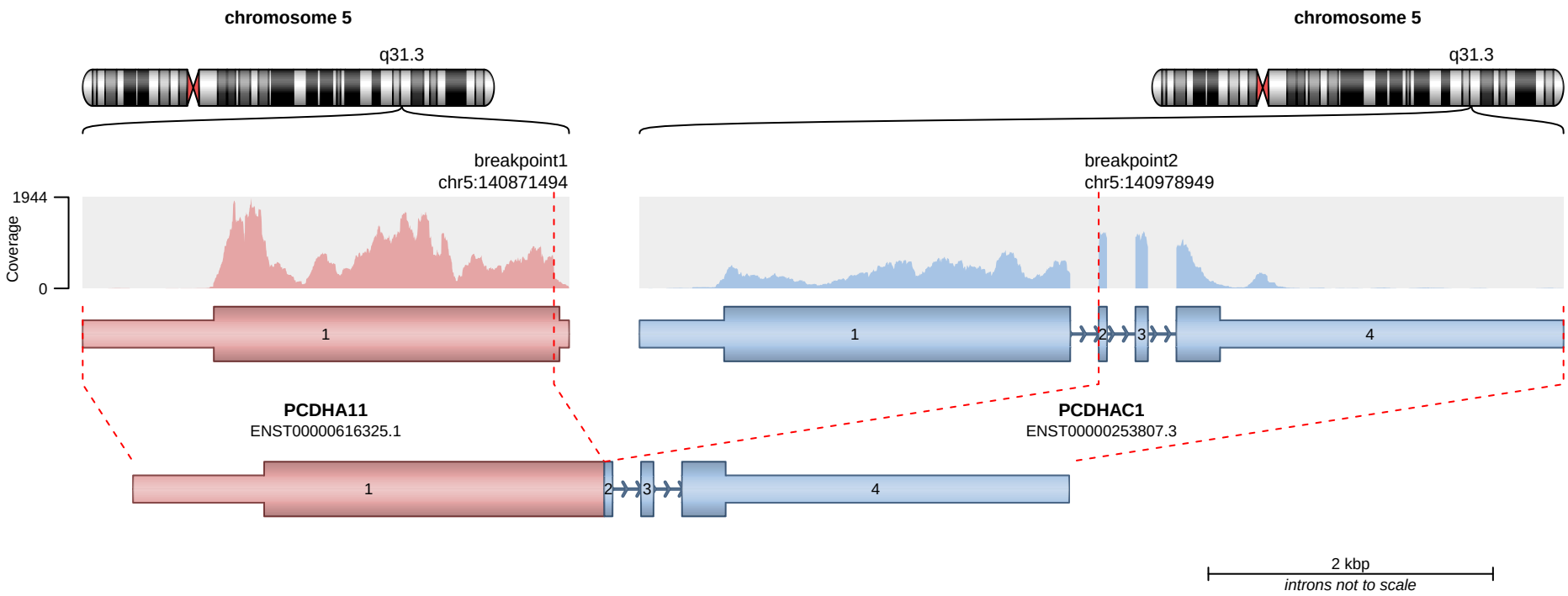


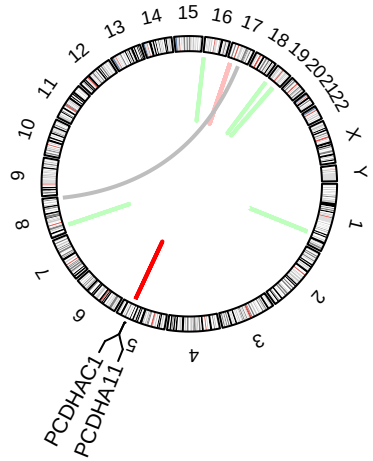


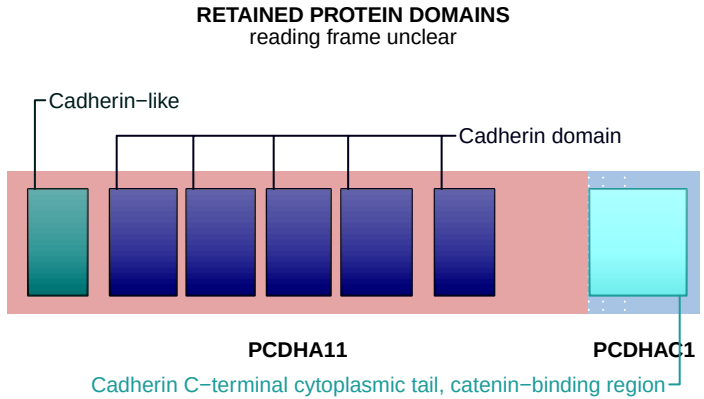
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 140
Discordant mates = 0



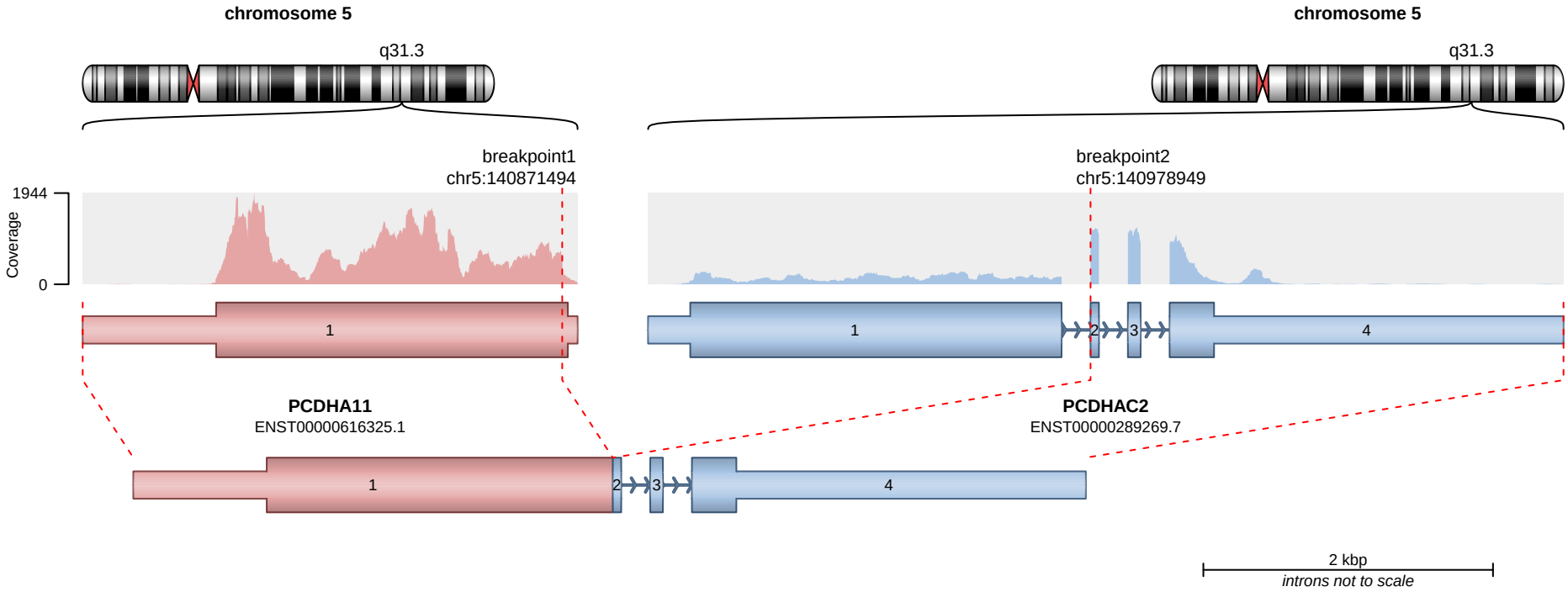


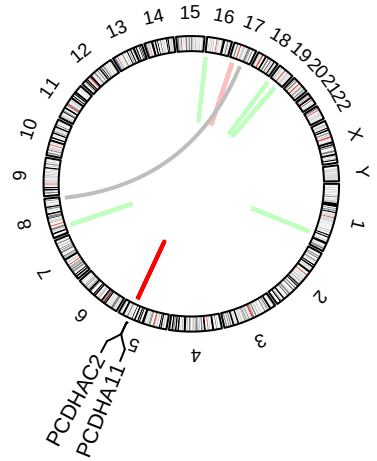


SUPPORTING READ COUNT

Split reads = 84
Discordant mates = 0

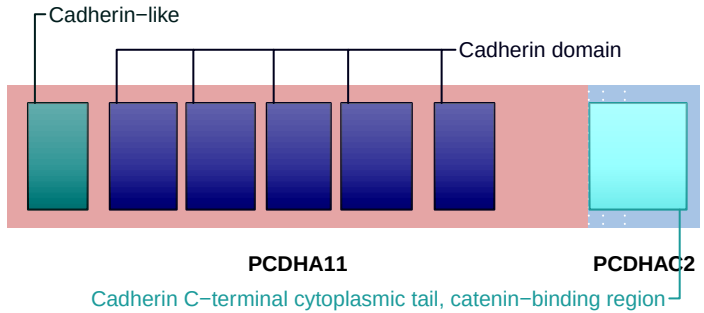
translocation
deletion
duplication
inversion





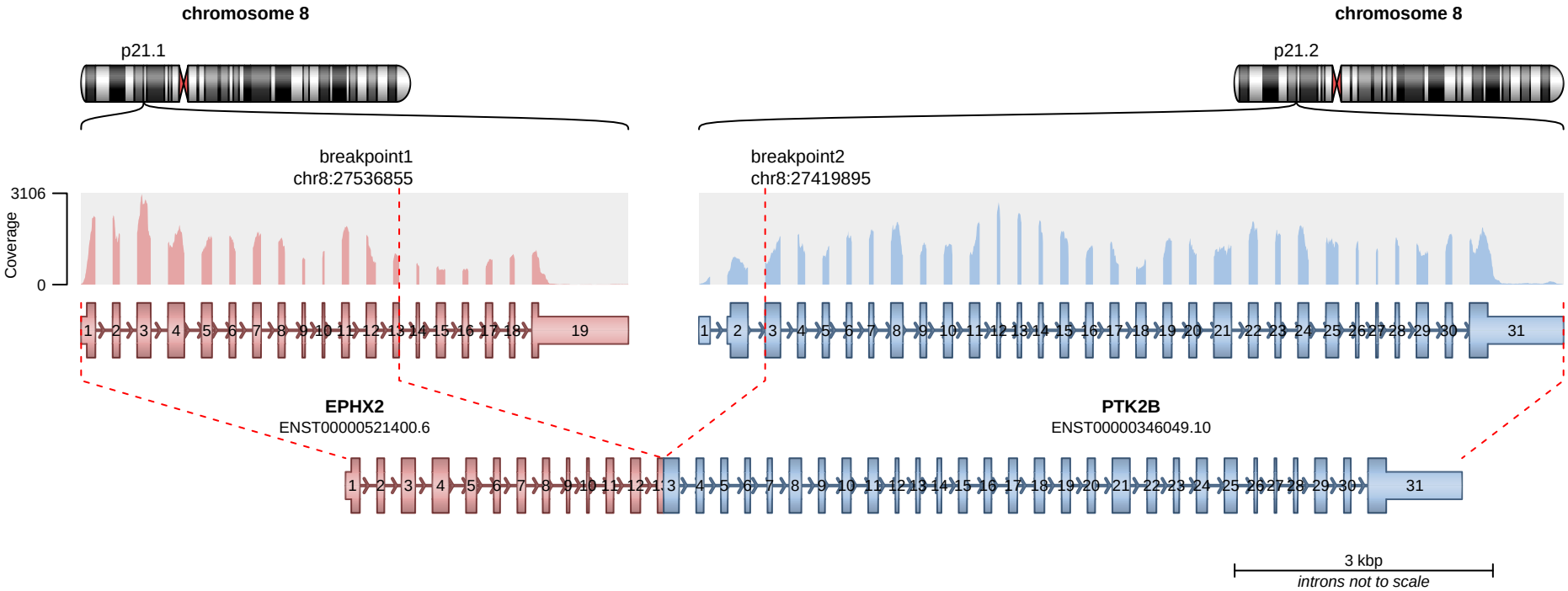
translocation
deletion
duplication
inversion

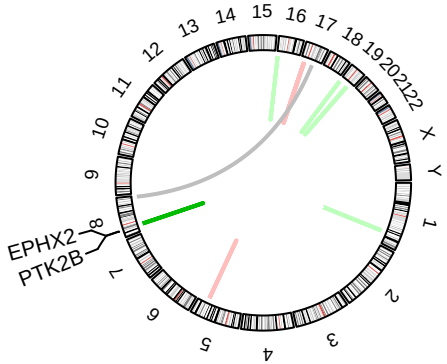
RETAINED PROTEIN DOMAINS
reading frame unclear

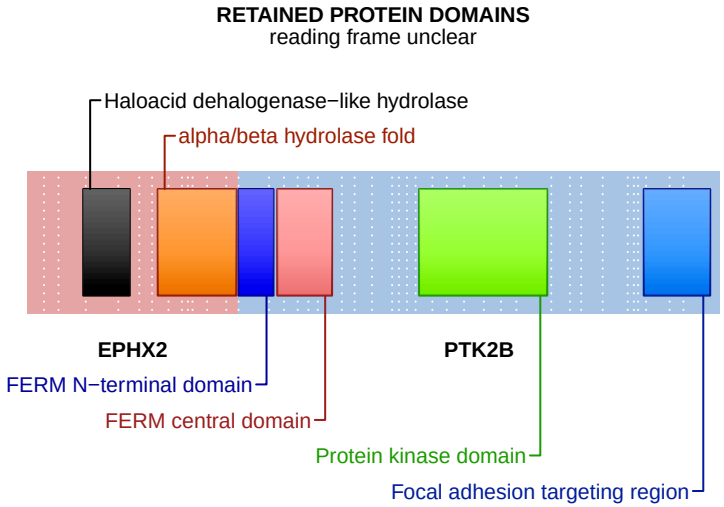


SUPPORTING READ COUNT

Split reads = 84
Discordant mates = 0

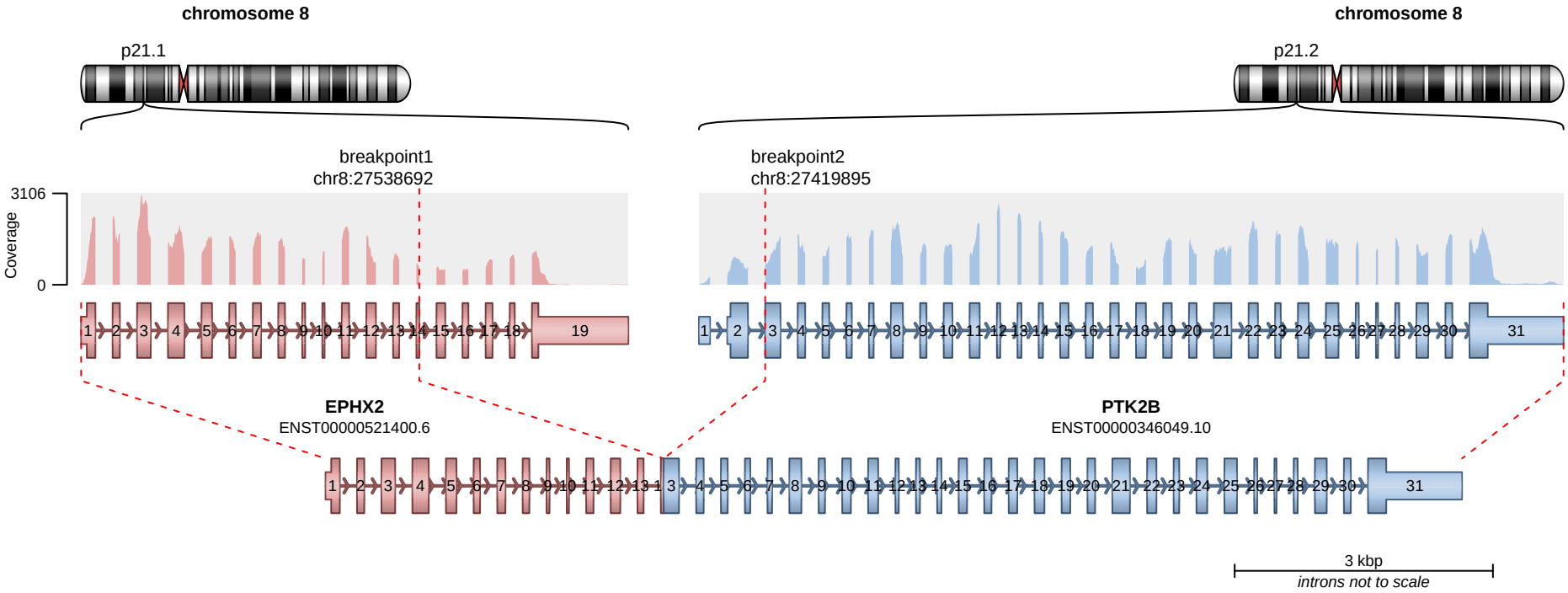


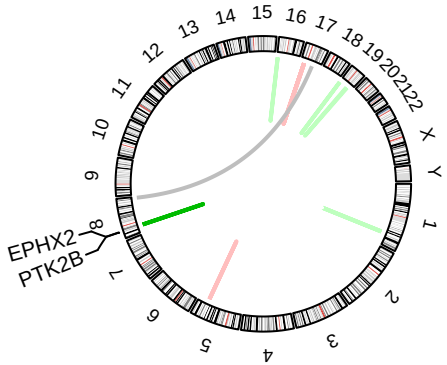




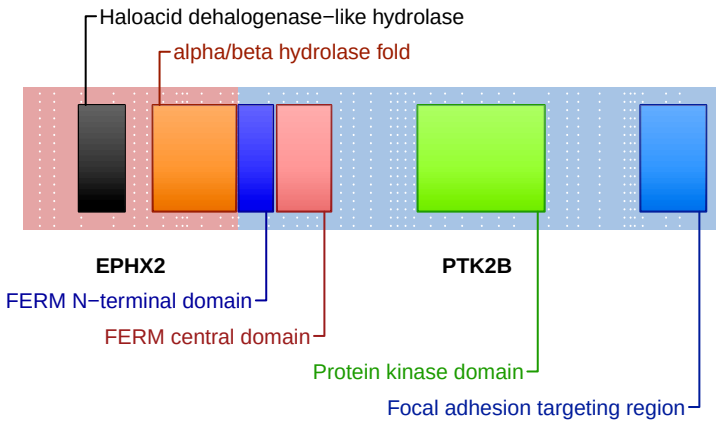
SUPPORTING READ COUNT

Split reads = 54
Discordant mates = 0



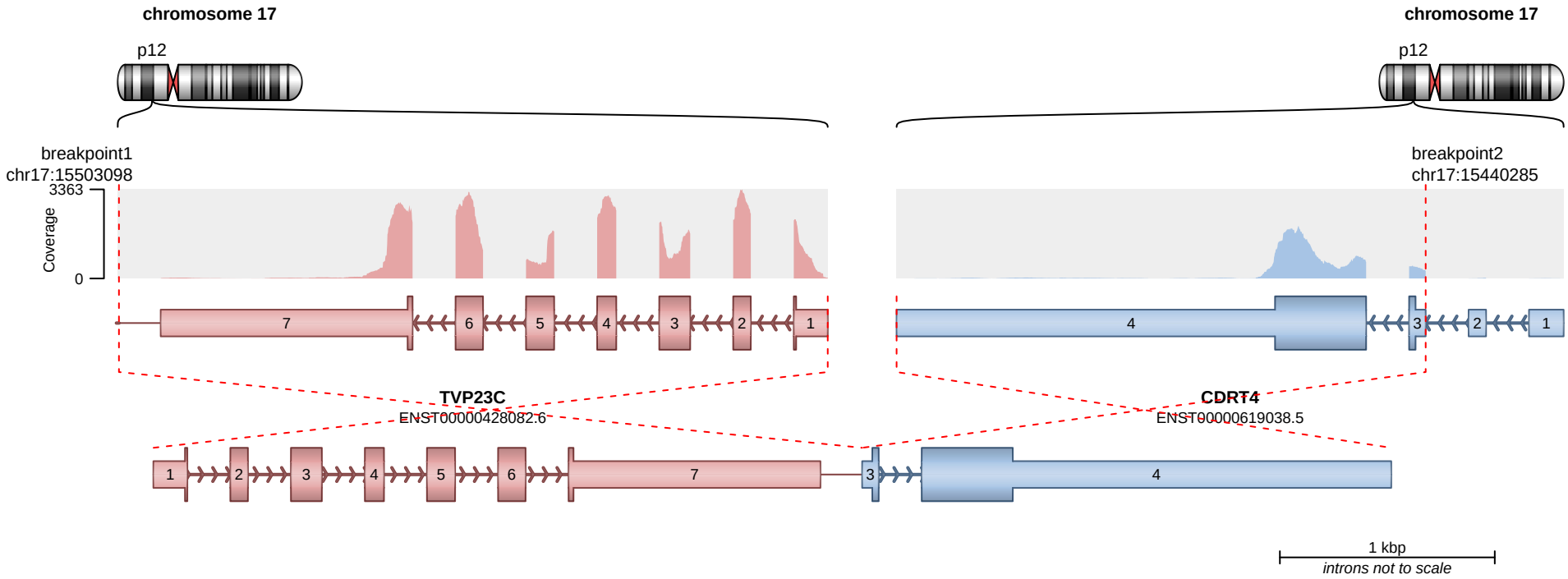


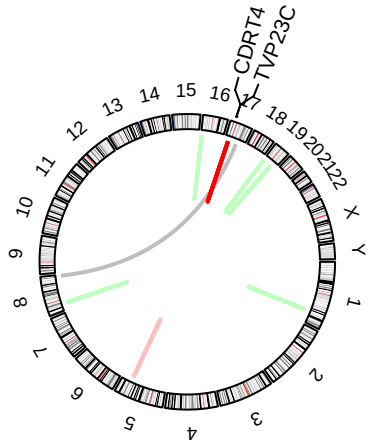
RETAINED PROTEIN DOMAINS
reading frame unclear



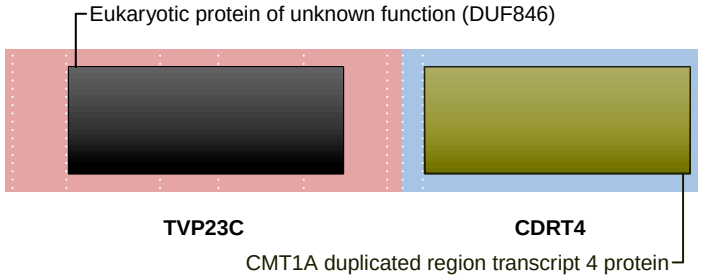
SUPPORTING READ COUNT

Split reads = 10
Discordant mates = 0





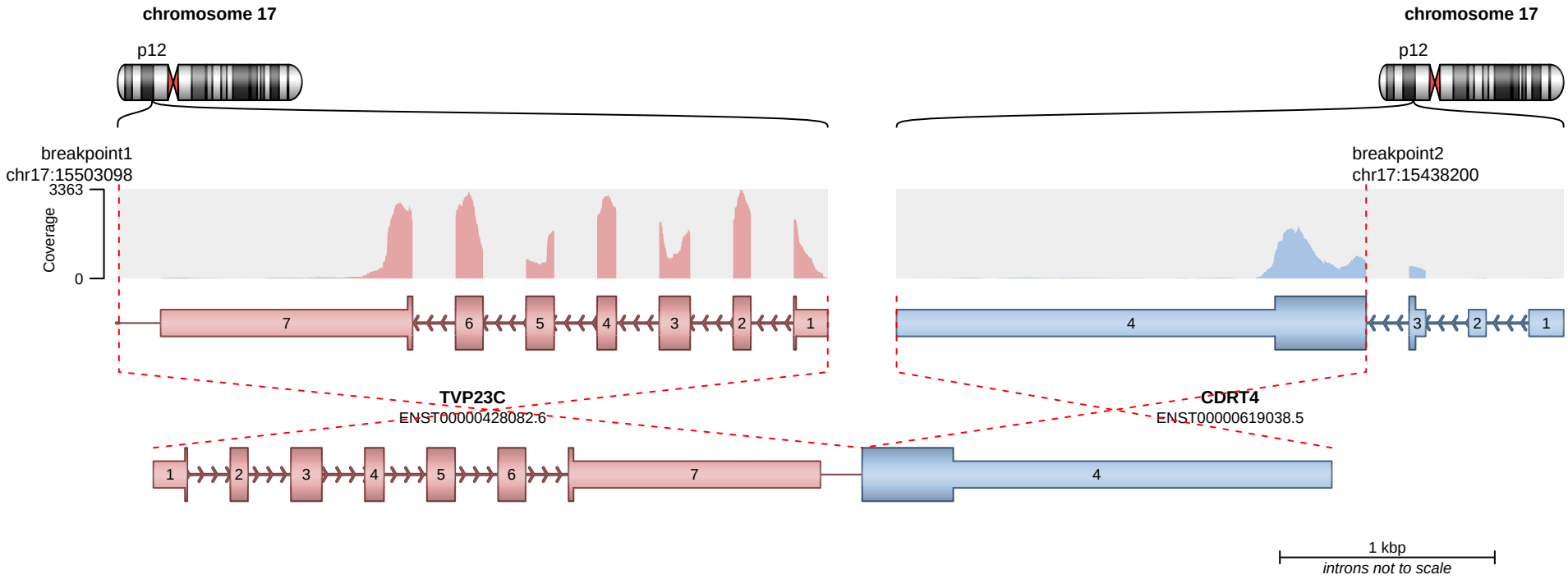
RETAINED PROTEIN DOMAINS
reading frame unclear

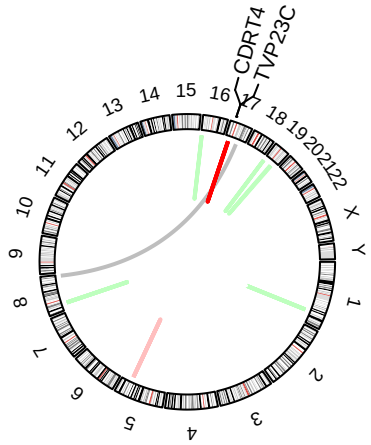


SUPPORTING READ COUNT

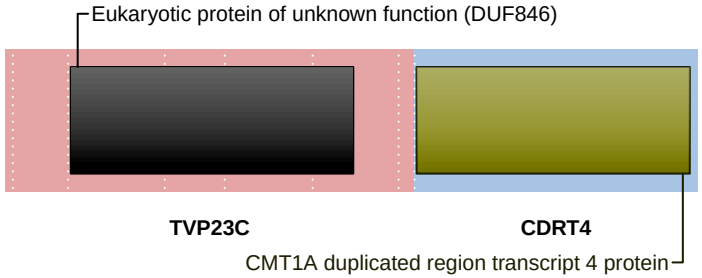
Split reads = 45
Discordant mates = 0

translocation
duplication
deletion
inversion





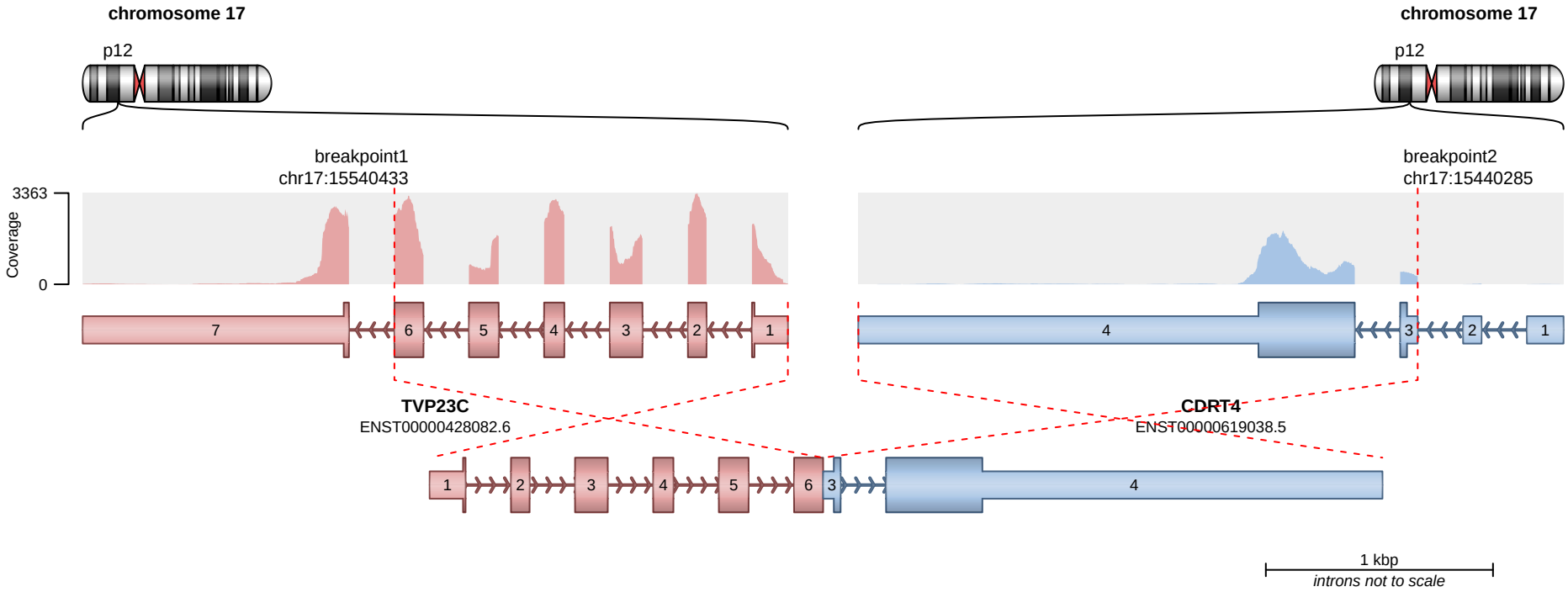
RETAINED PROTEIN DOMAINS
reading frame unclear

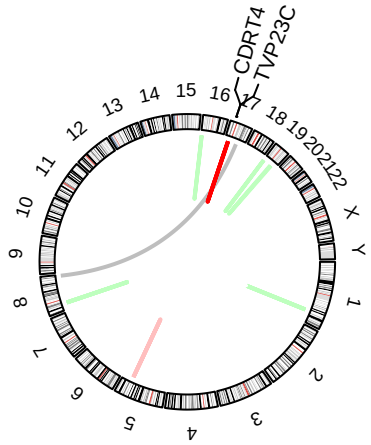


SUPPORTING READ COUNT

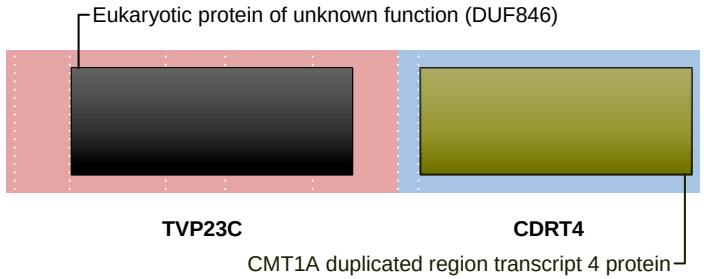
Split reads = 43
Discordant mates = 0

translocation
duplication
deletion
inversion



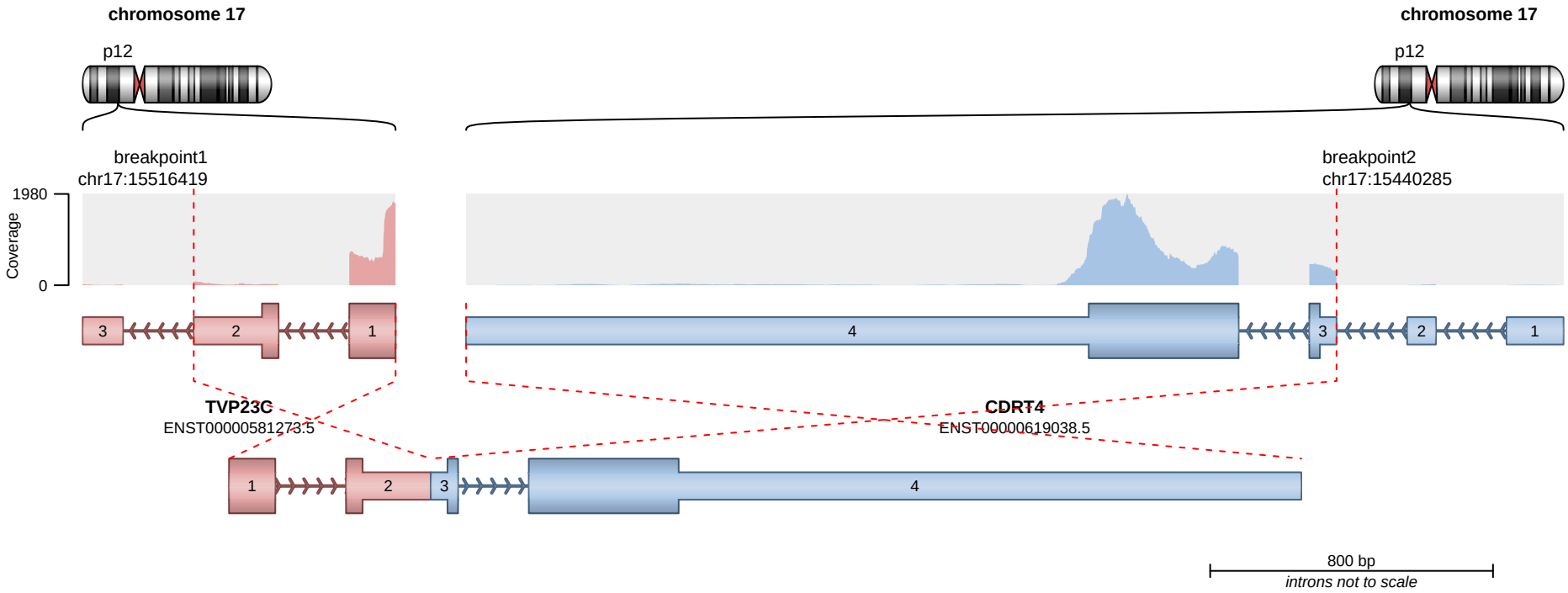


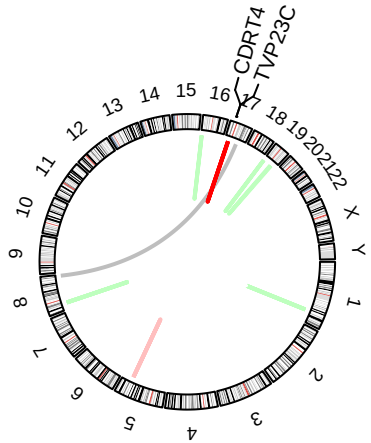
RETAINED PROTEIN DOMAINS
reading frame unclear



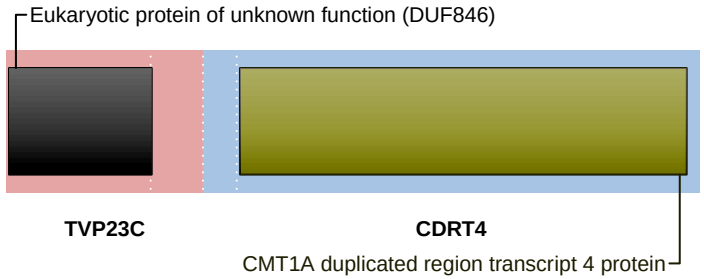
SUPPORTING READ COUNT

Split reads = 27
Discordant mates = 0





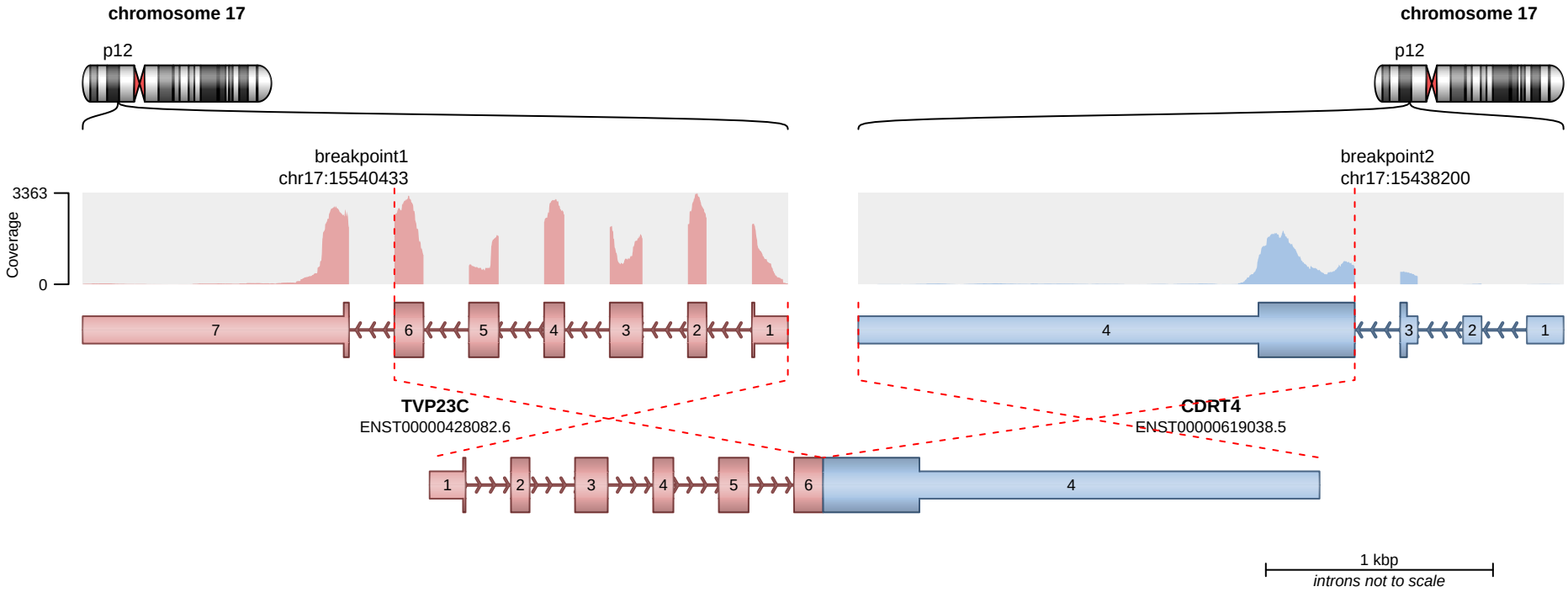
RETAINED PROTEIN DOMAINS
reading frame unclear

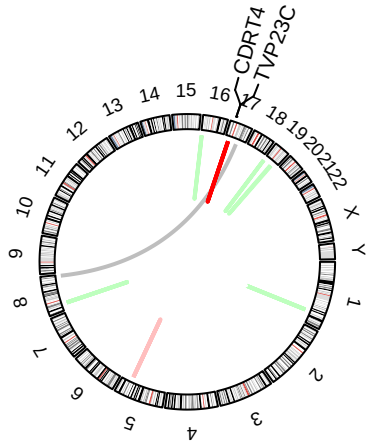


SUPPORTING READ COUNT

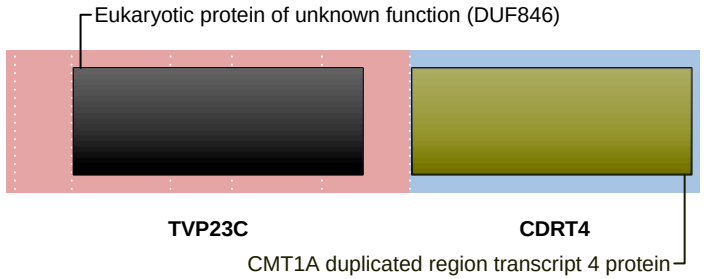
Split reads = 2
Discordant mates = 0

translocation
deletion
duplication
inversion





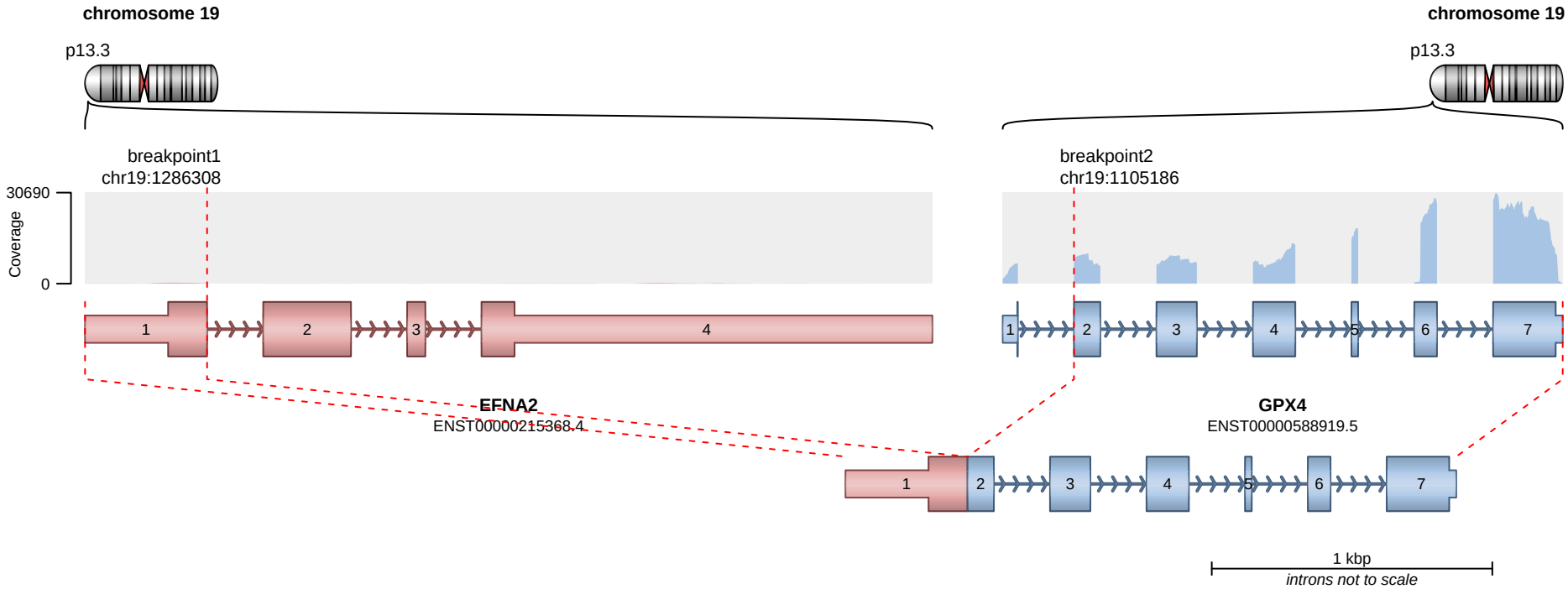
RETAINED PROTEIN DOMAINS
reading frame unclear

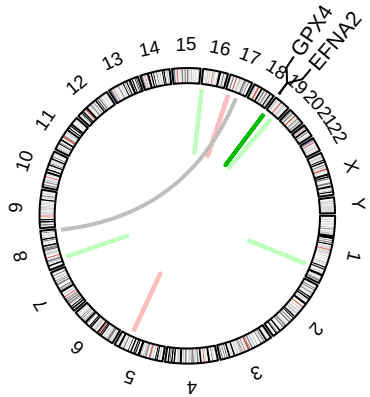


SUPPORTING READ COUNT

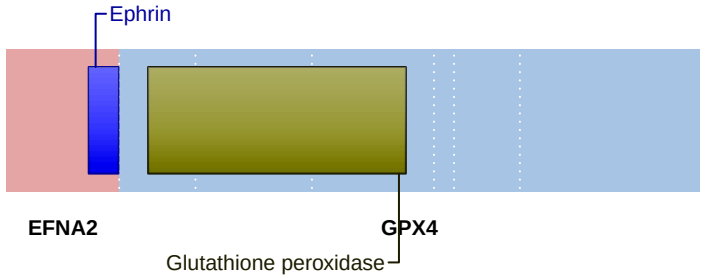
Split reads = 1
Discordant mates = 0

translocation
duplication
deletion
inversion



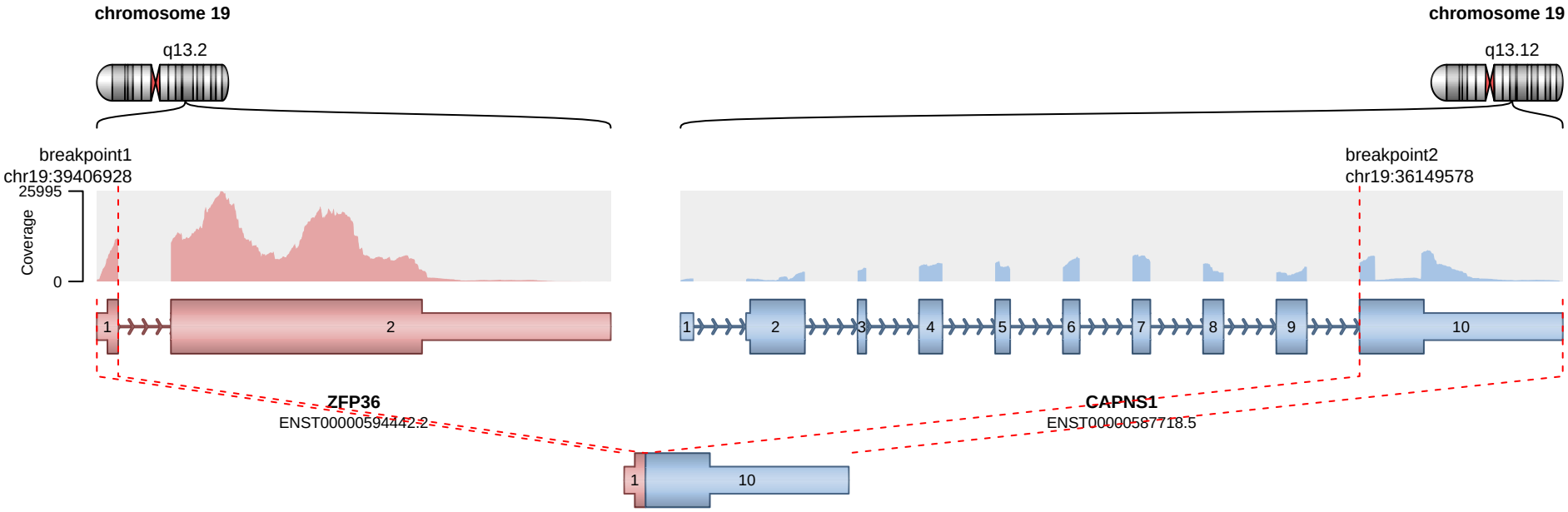


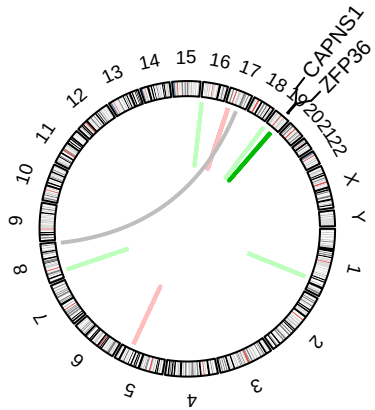
RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 37
Discordant mates = 0



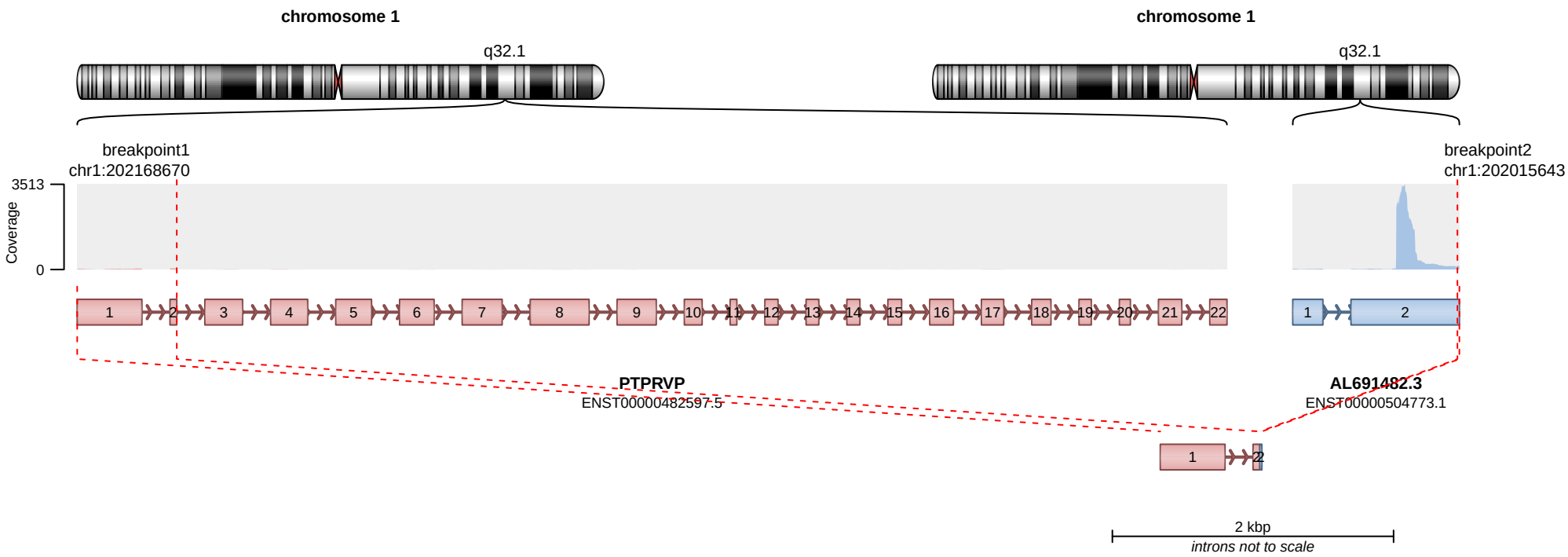


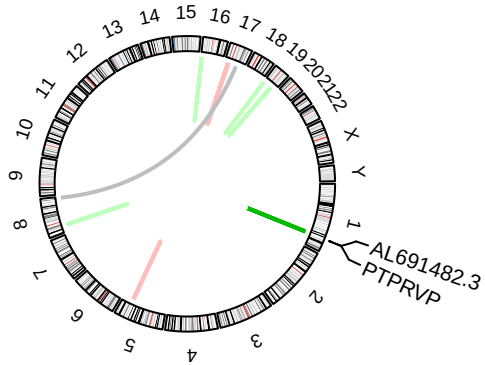
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 23
Discordant mates = 0

translocation
deletion
duplication
inversion



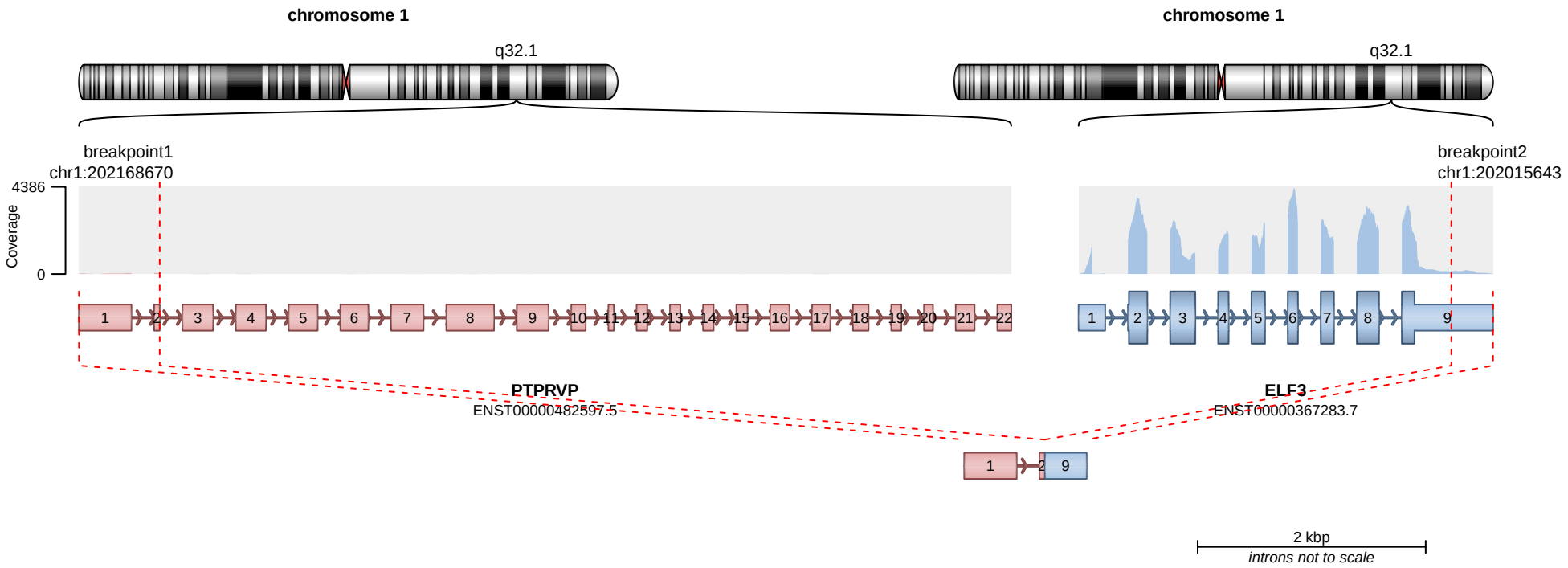


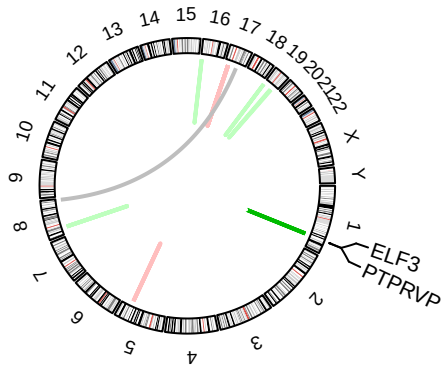
Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 15
Discordant mates = 0

translocation
deletion
duplication
inversion



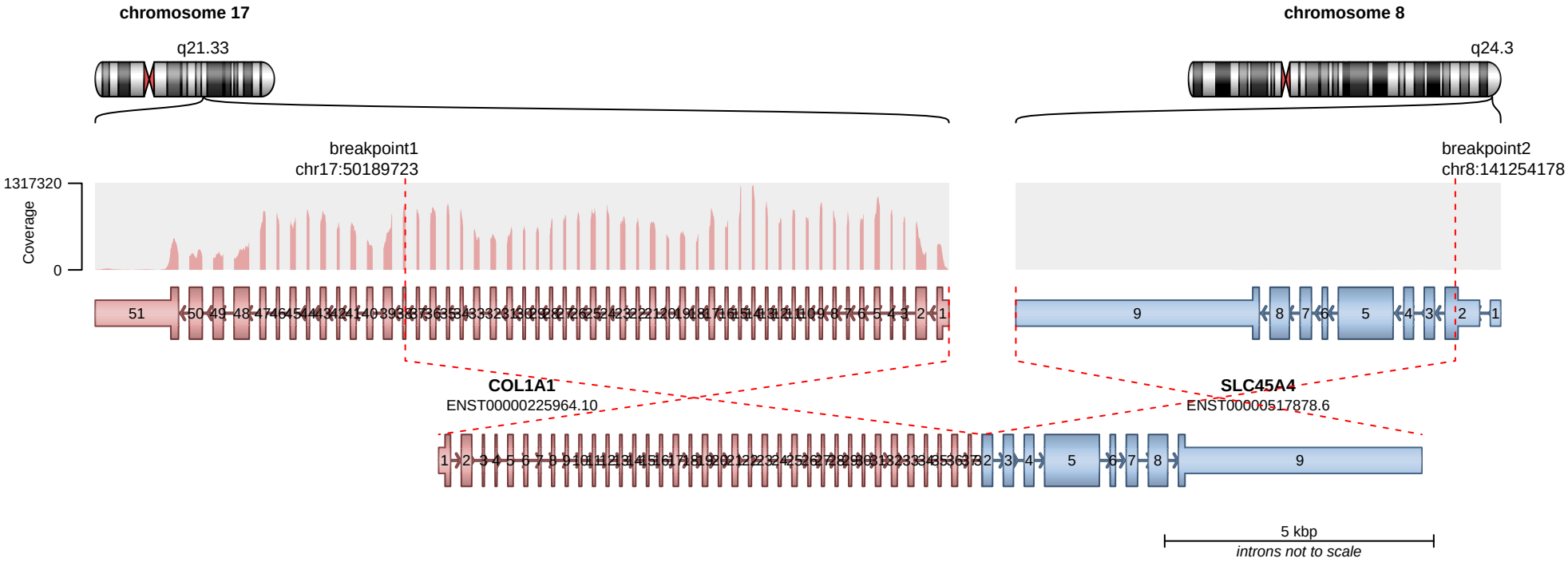


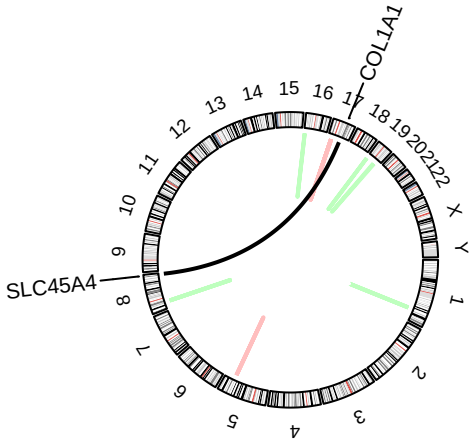
No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

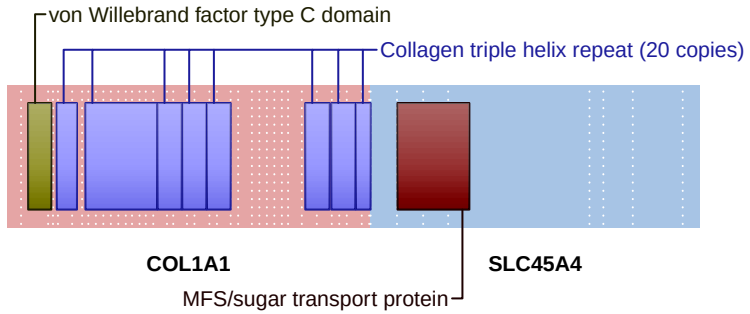
Split reads = 15
Discordant mates = 0

translocation
deletion
duplication
inversion





RETAINED PROTEIN DOMAINS
reading frame unclear



SUPPORTING READ COUNT

Split reads = 6
Discordant mates = 0

translocation
deletion
duplication
inversion