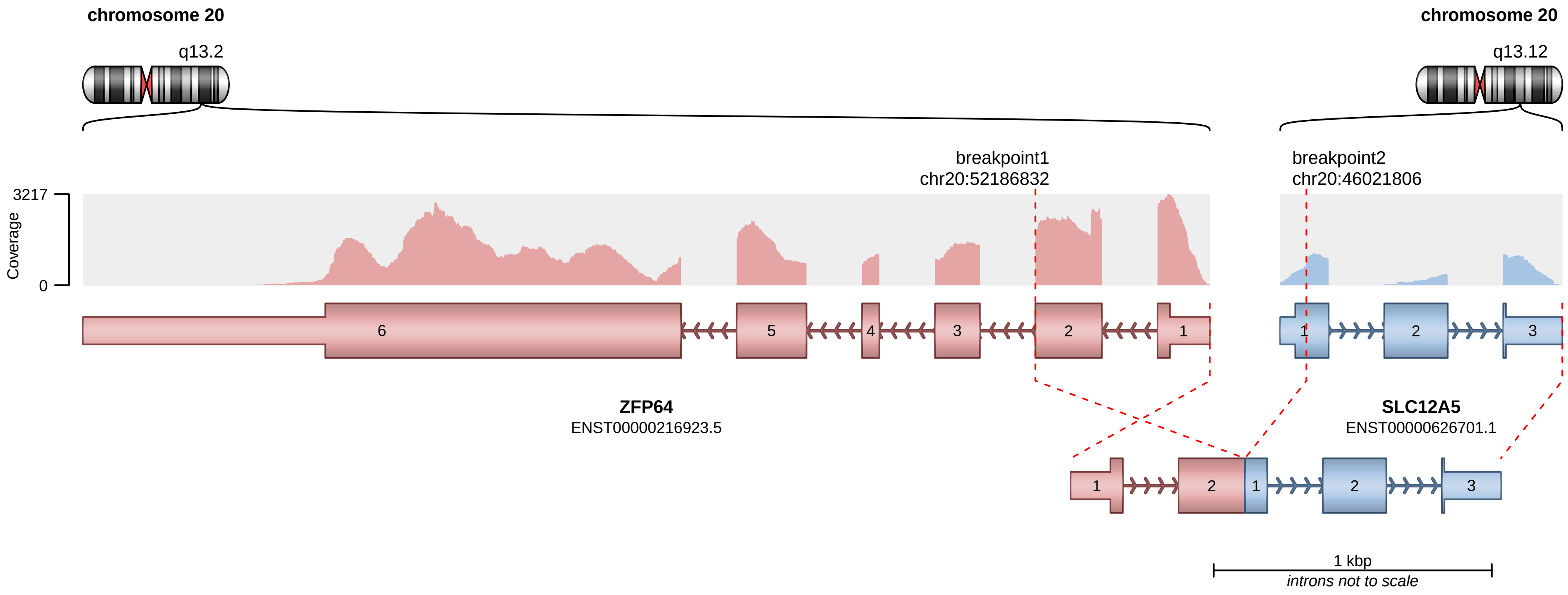
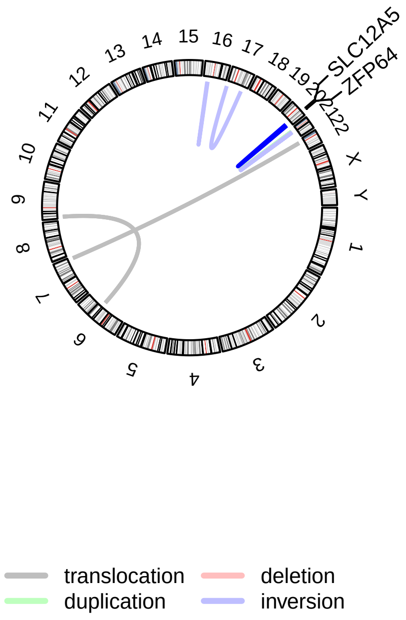
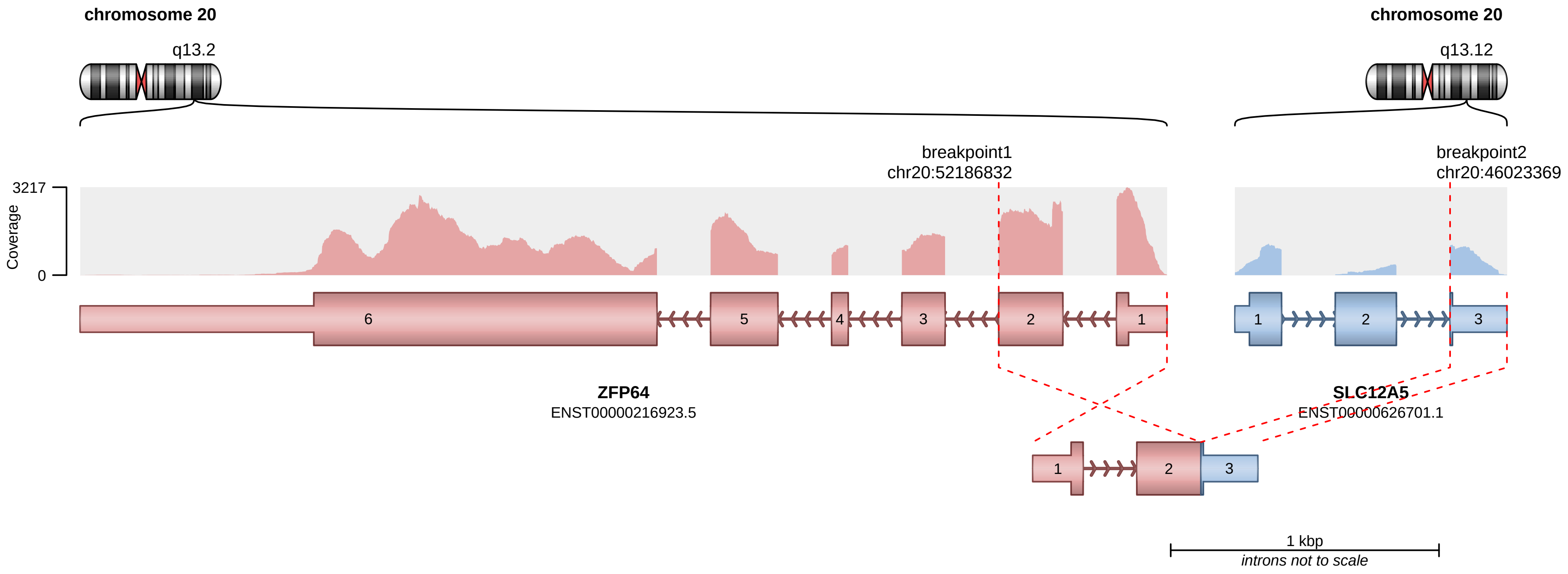


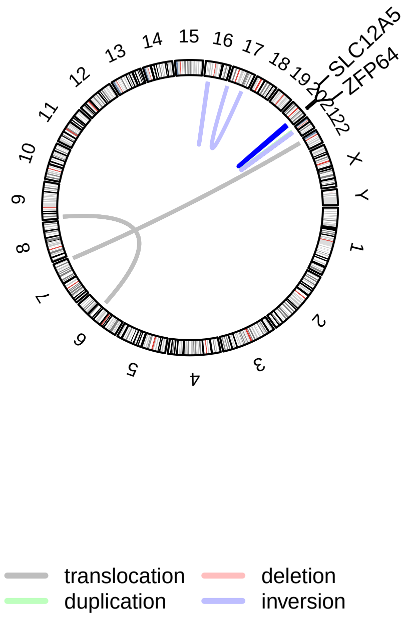


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 85
Discordant mates = 0

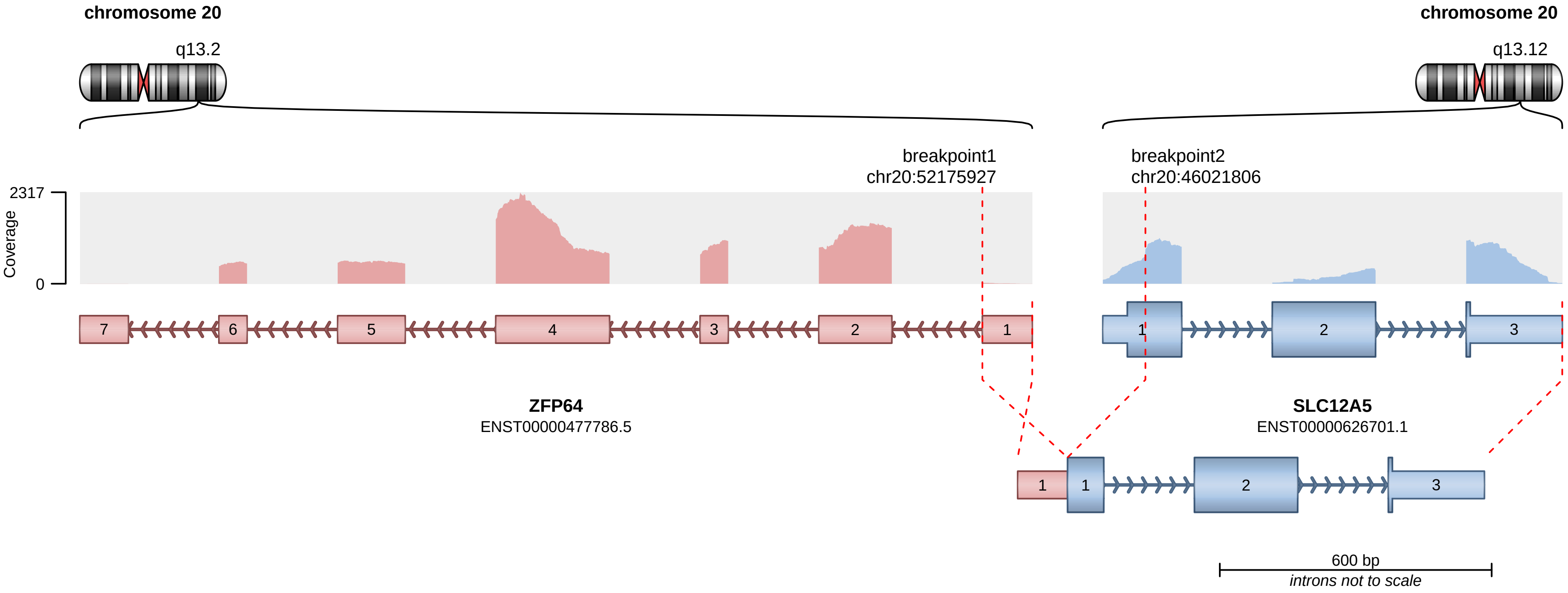


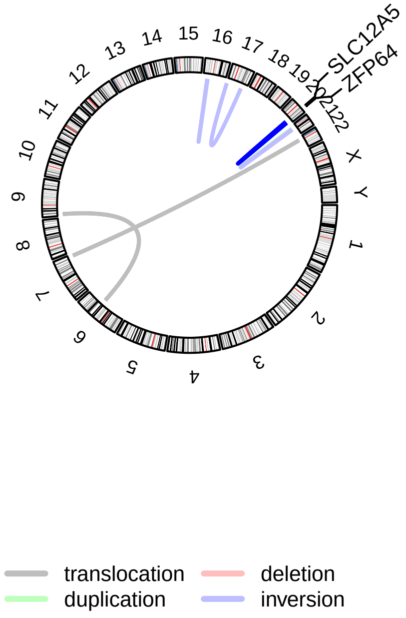


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 18
Discordant mates = 0

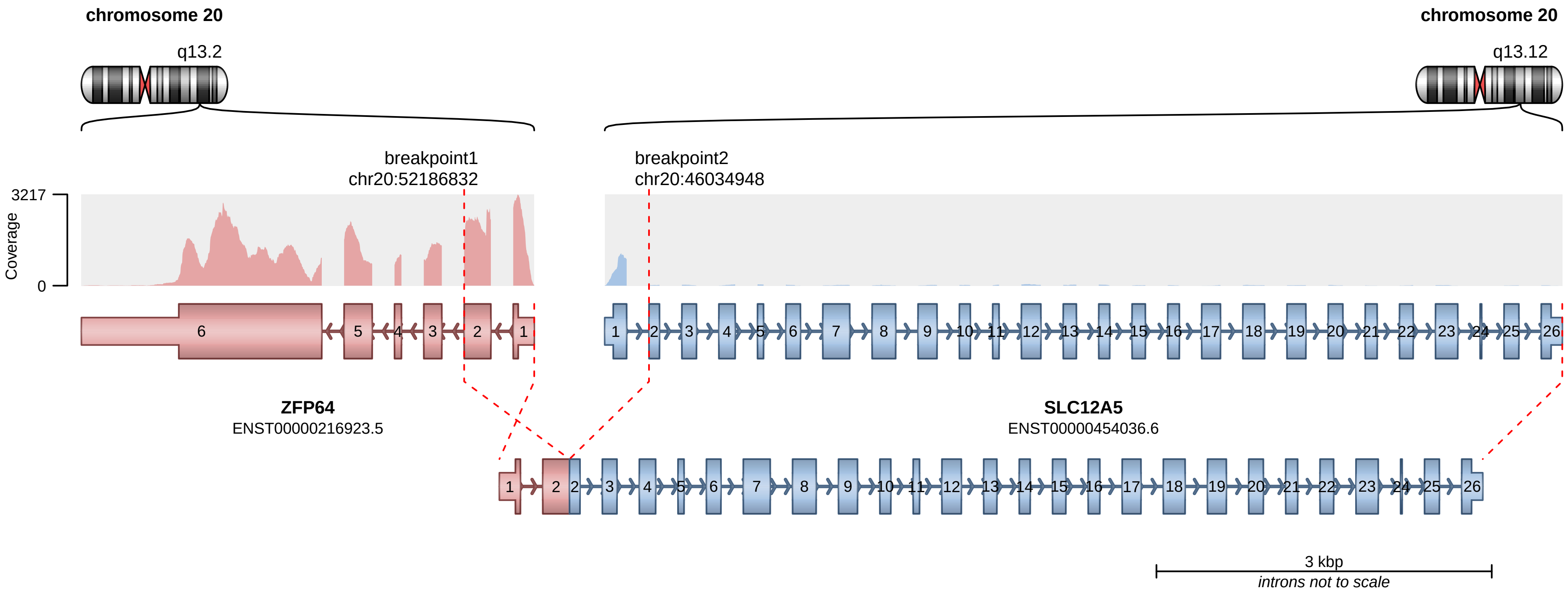


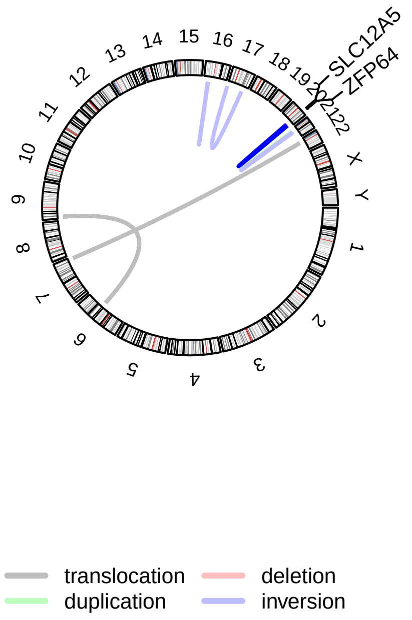


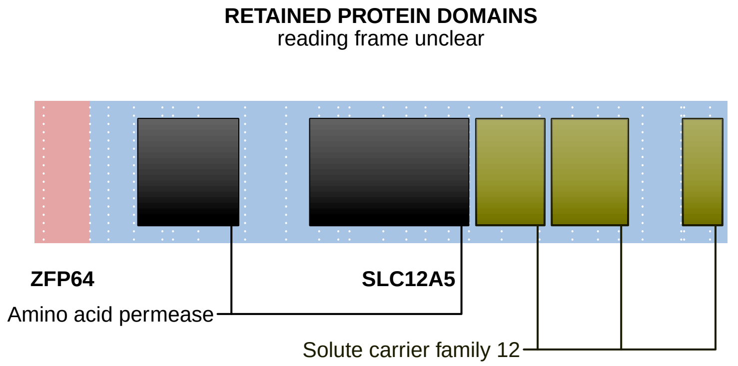
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 0

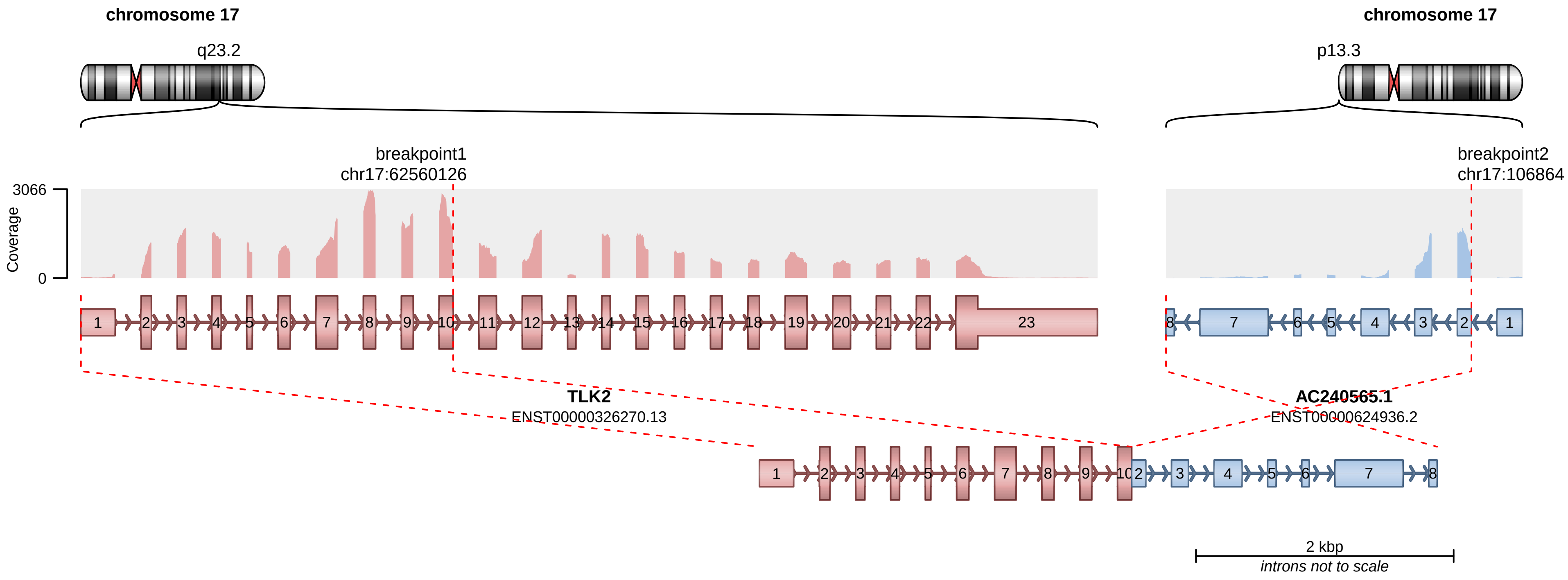


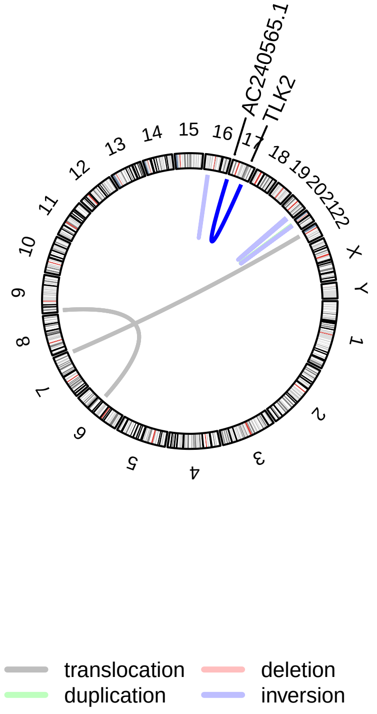




SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 0

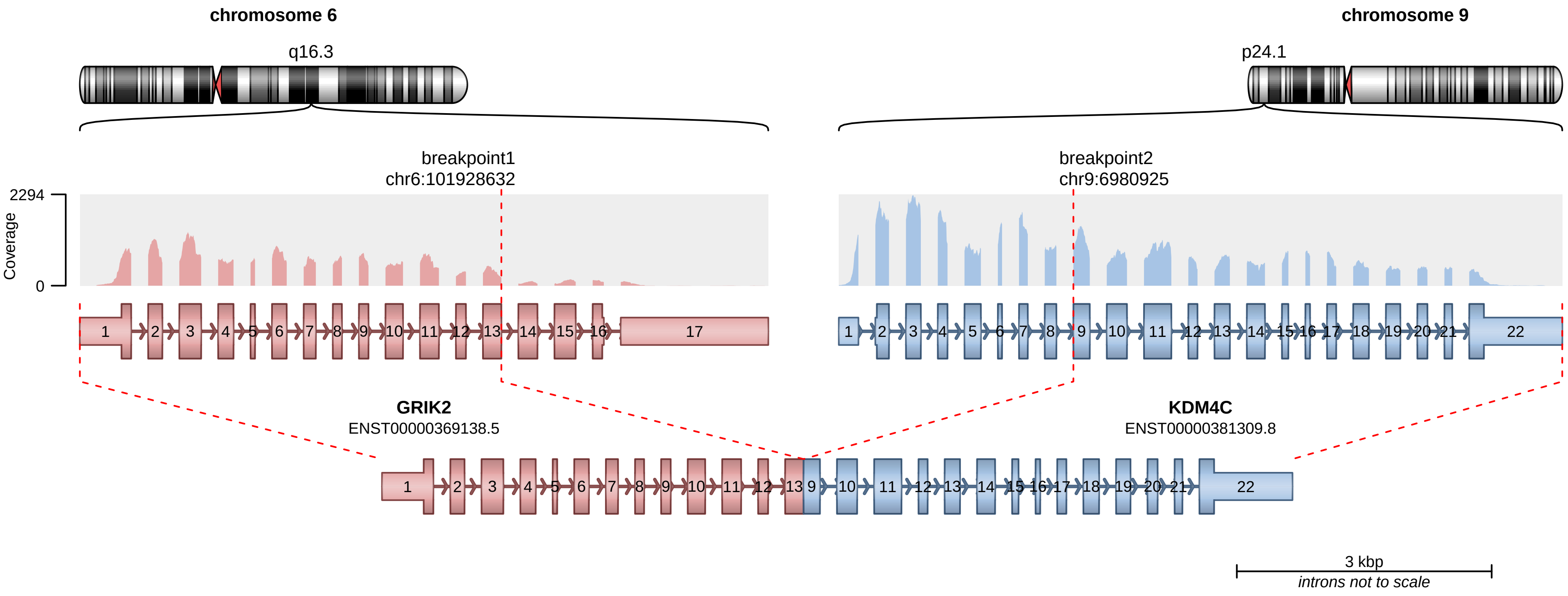


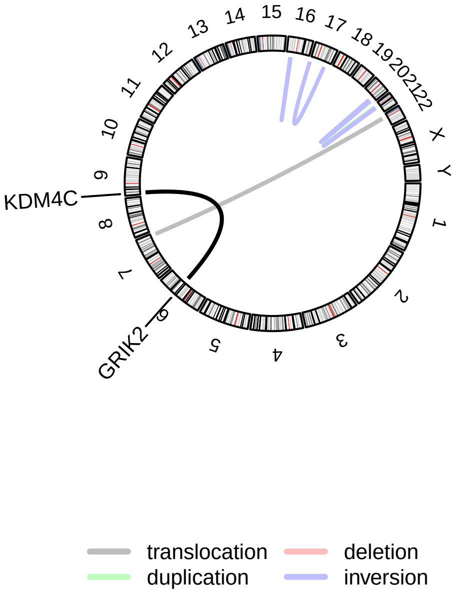


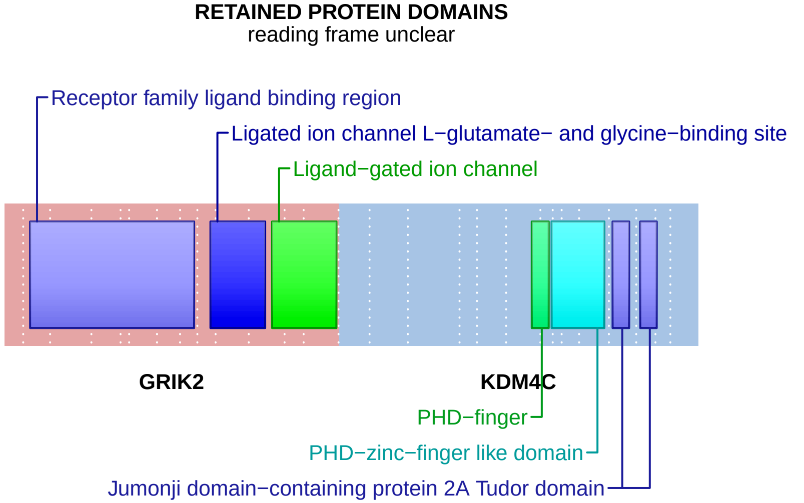
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 81
Discordant mates = 0

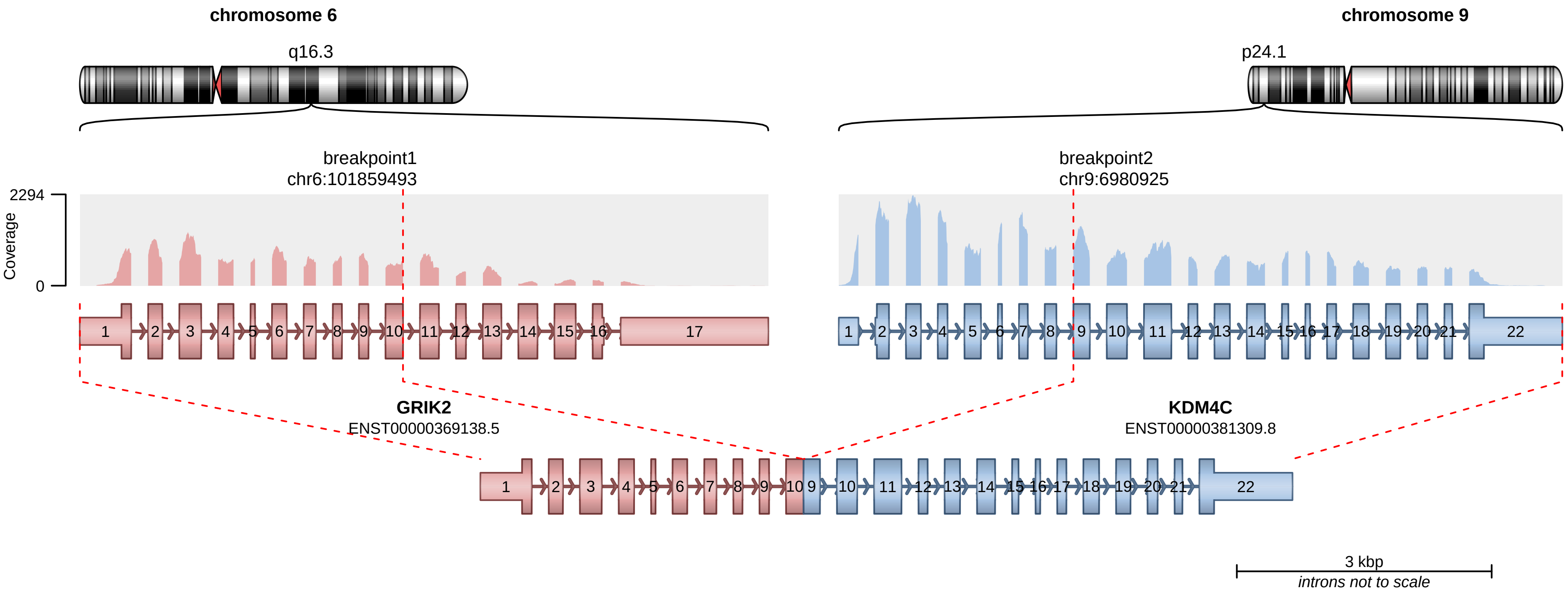


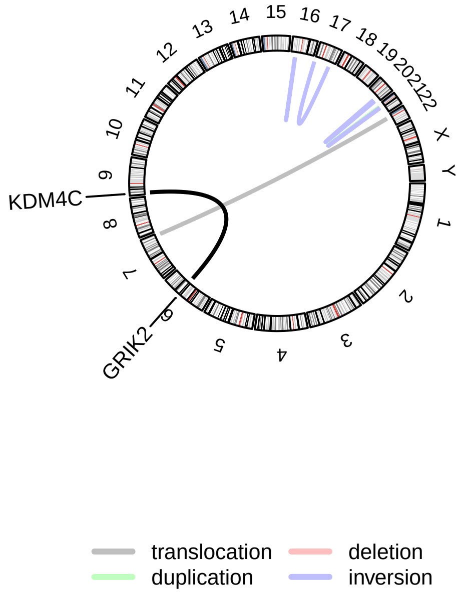


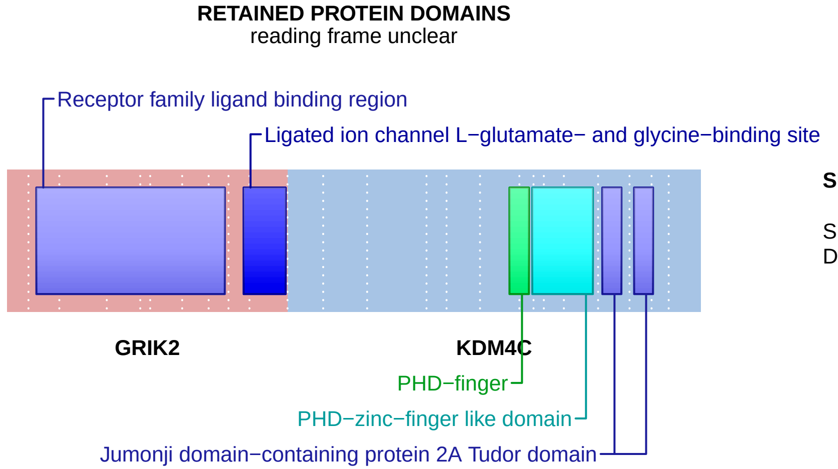


SUPPORTING READ COUNT

Split reads = 70
Discordant mates = 0

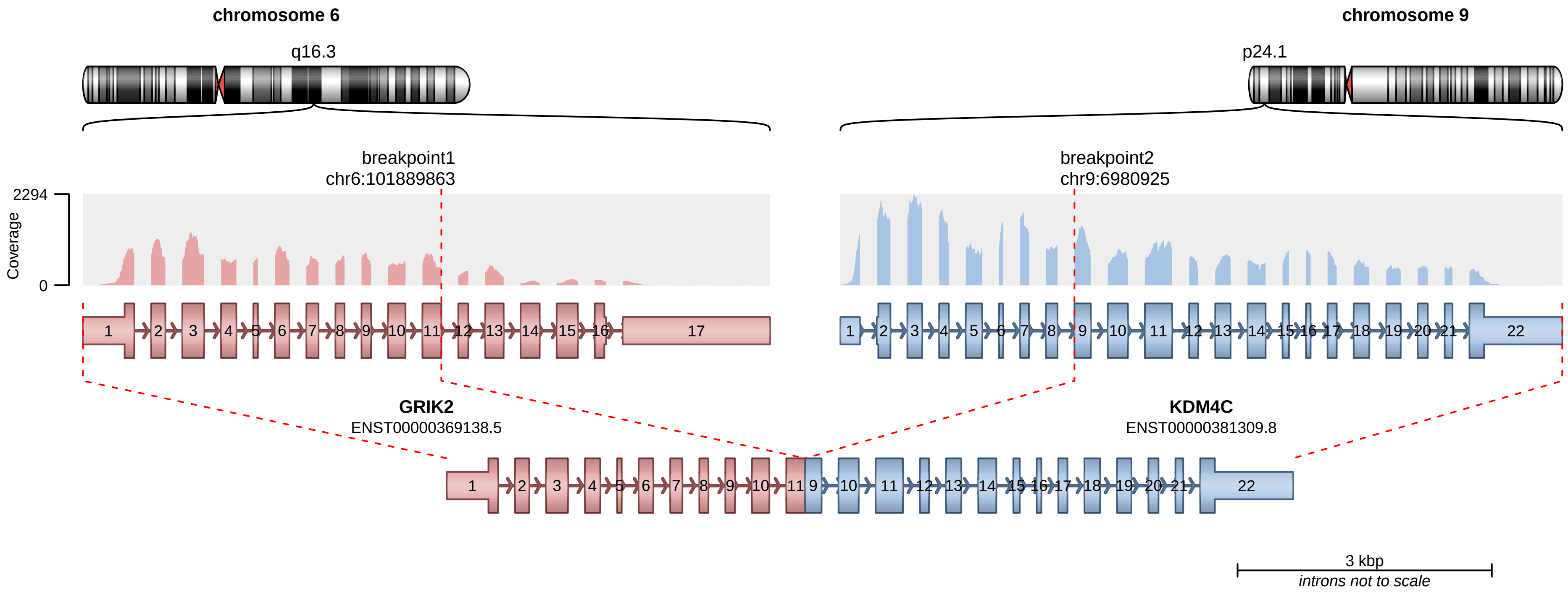


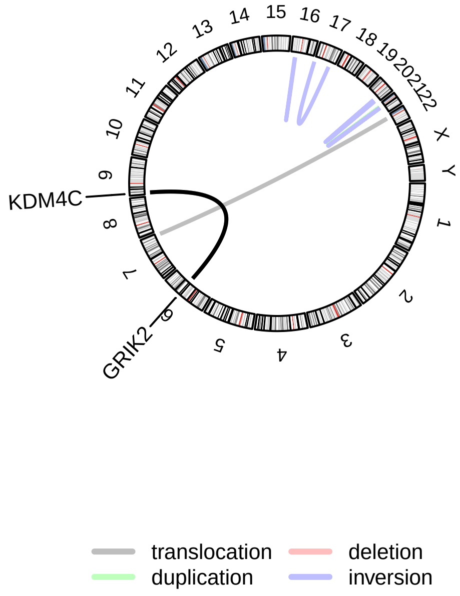


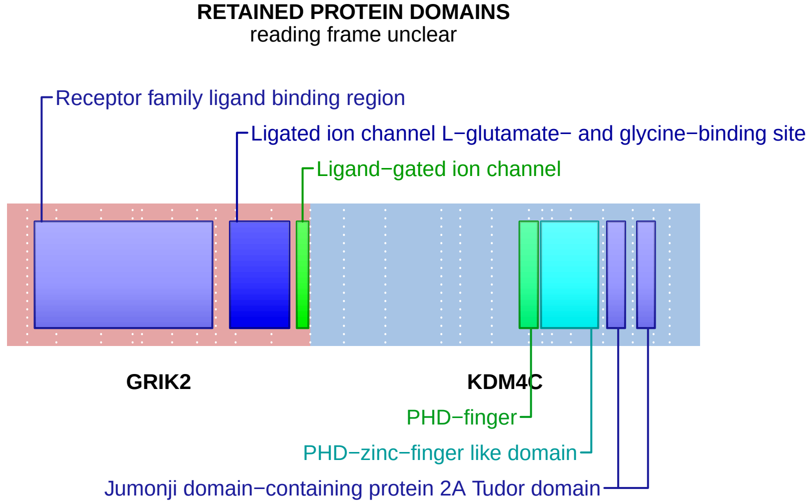


SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

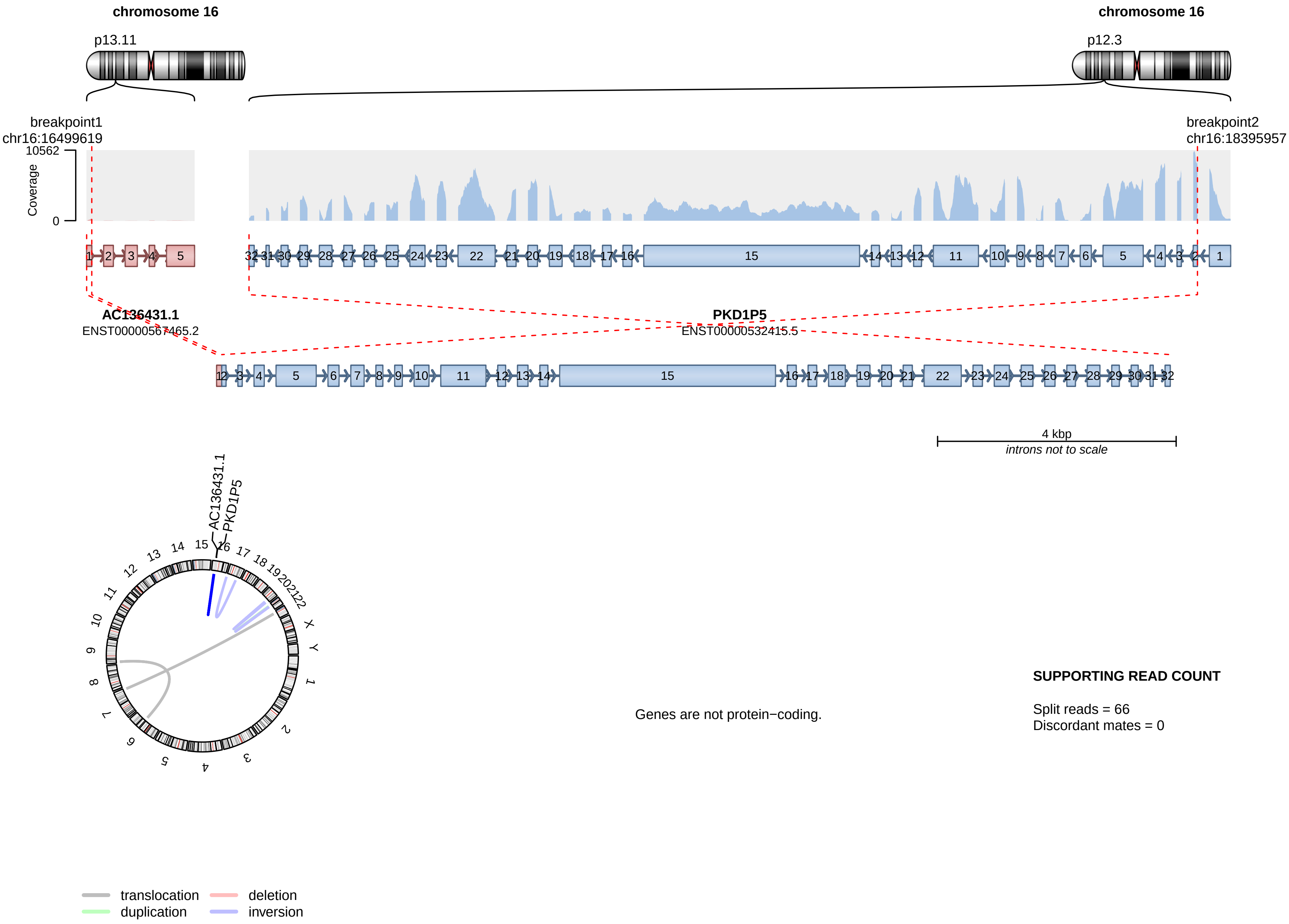






SUPPORTING READ COUNT

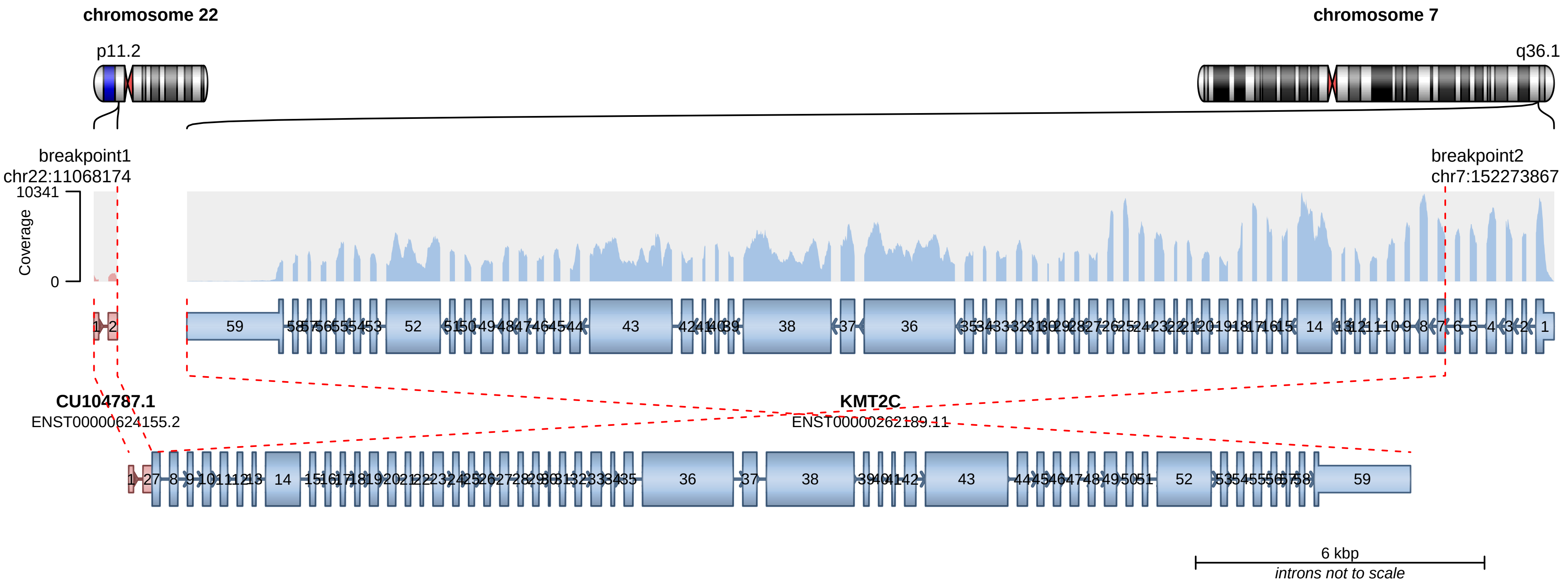
Split reads = 1
Discordant mates = 0

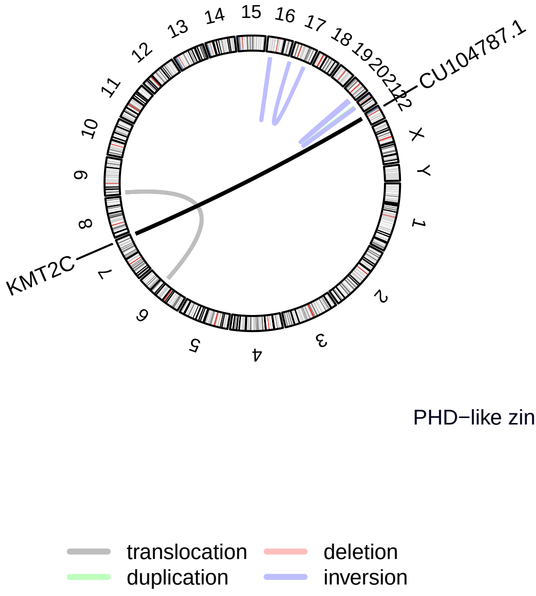


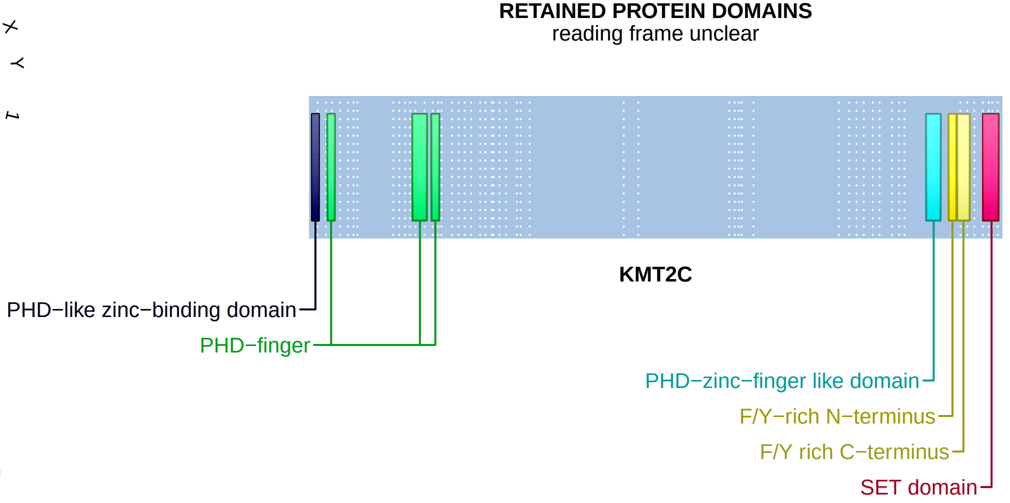
Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 66
Discordant mates = 0

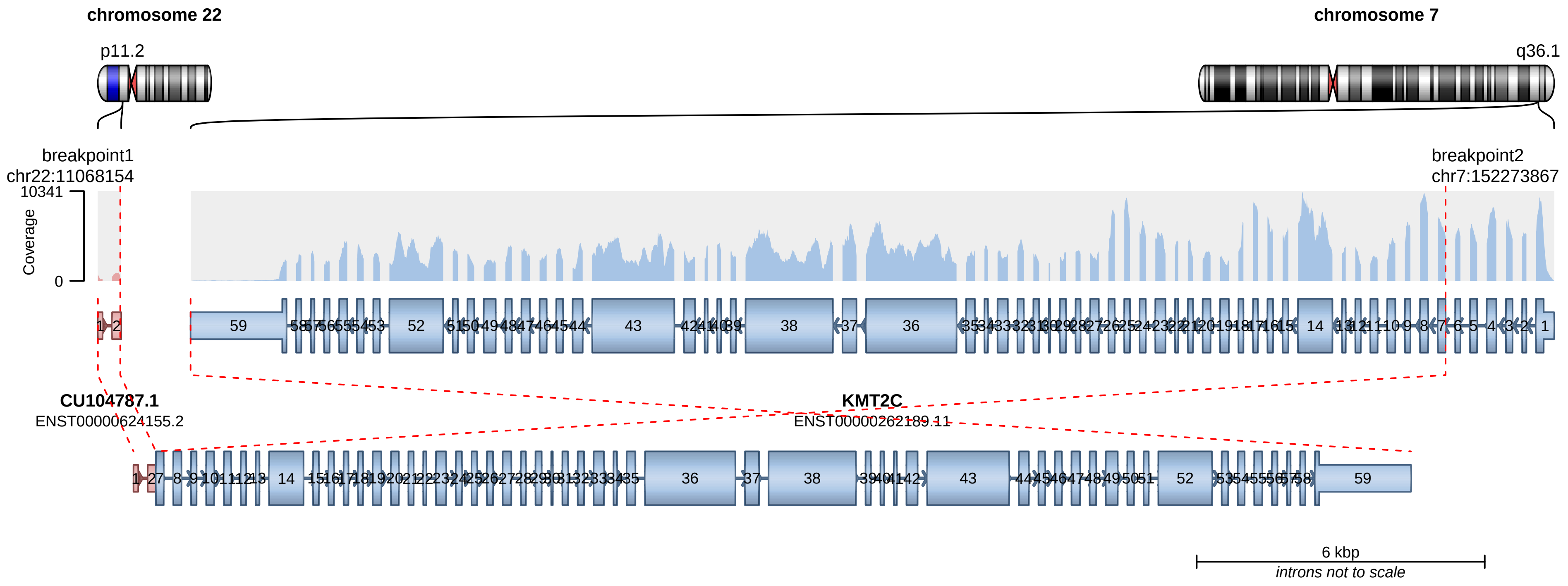


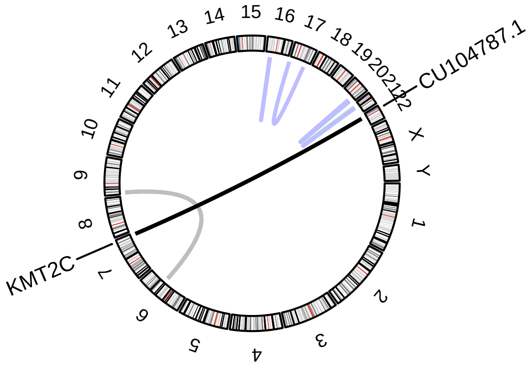




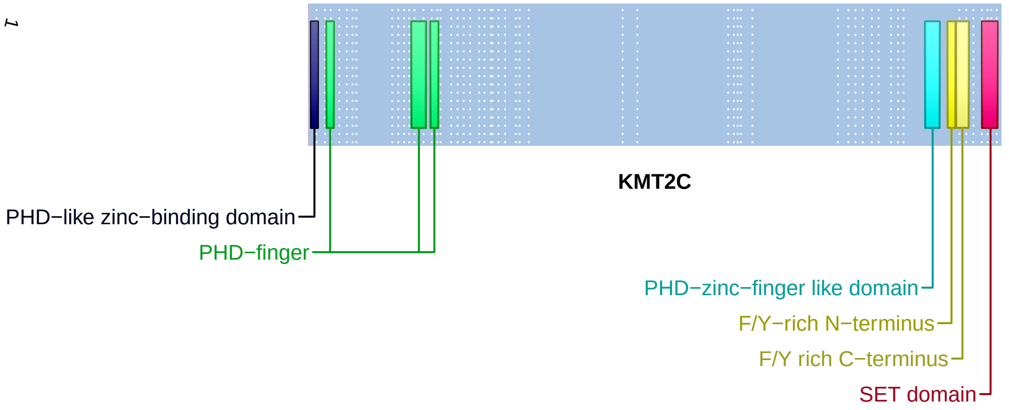
SUPPORTING READ COUNT

Split reads = 42
Discordant mates = 1



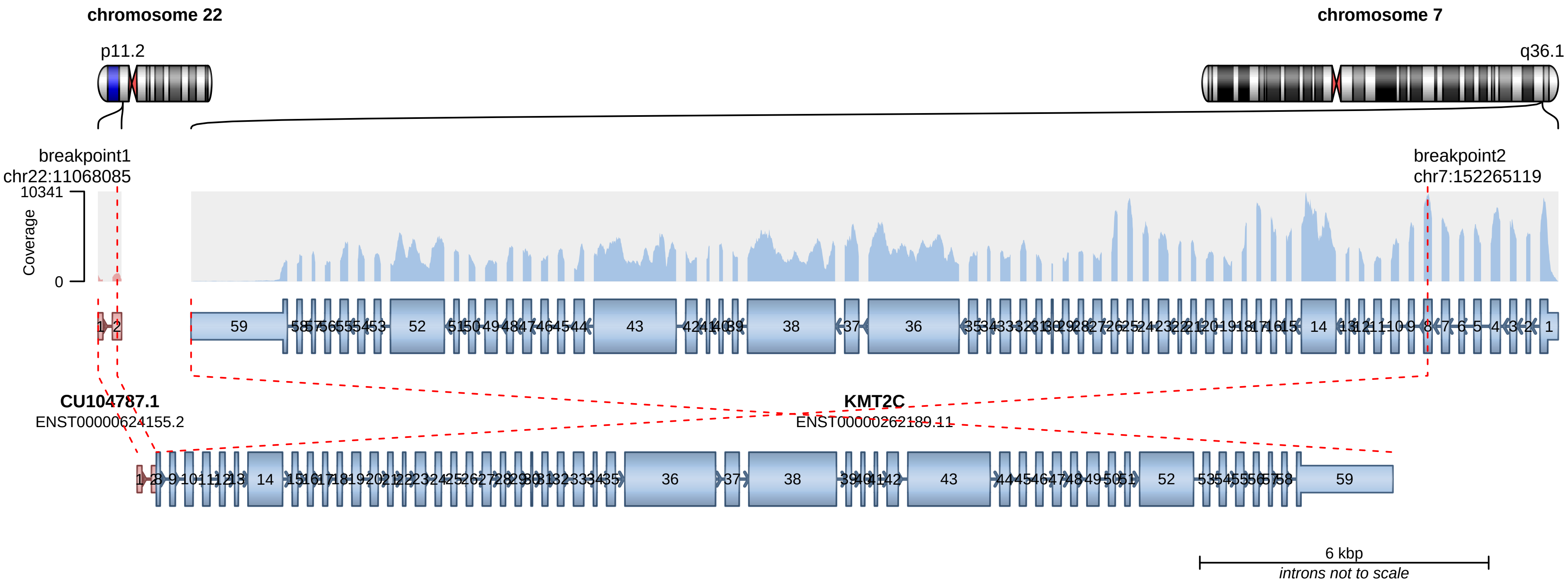


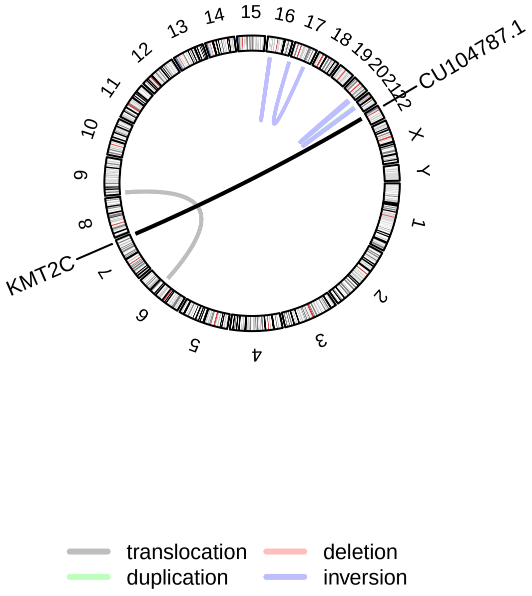
RETAINED PROTEIN DOMAINS
reading frame unclear

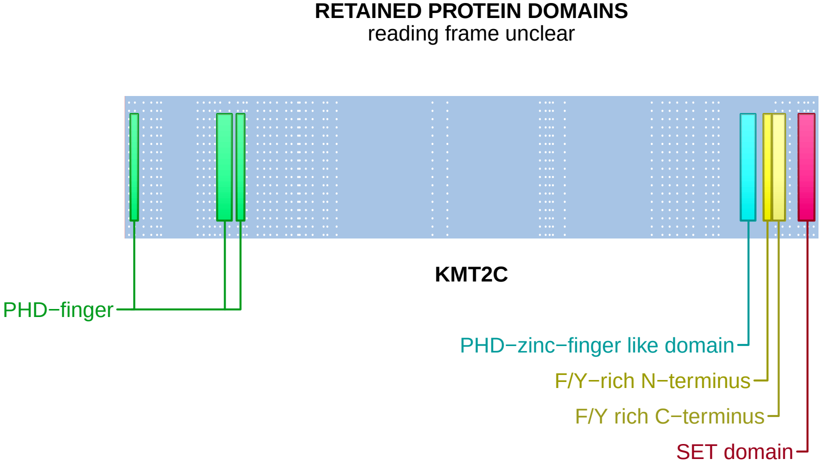


SUPPORTING READ COUNT

Split reads = 24
Discordant mates = 1

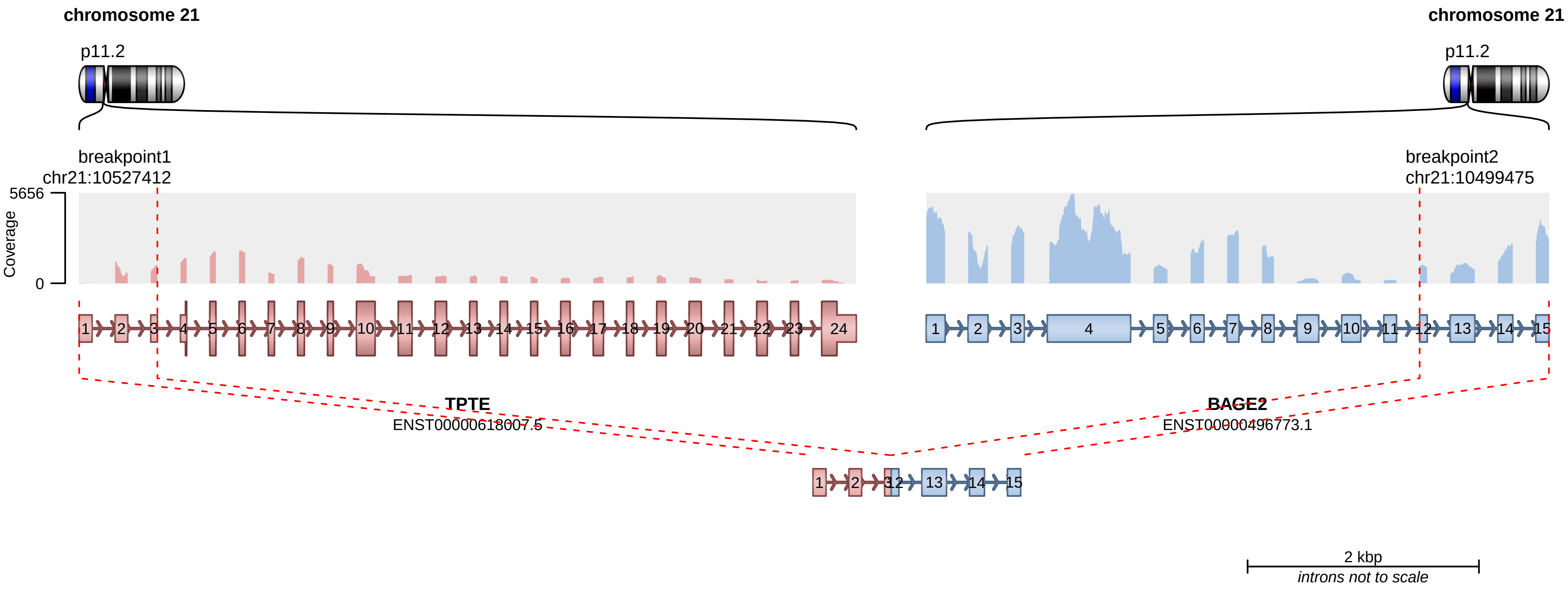


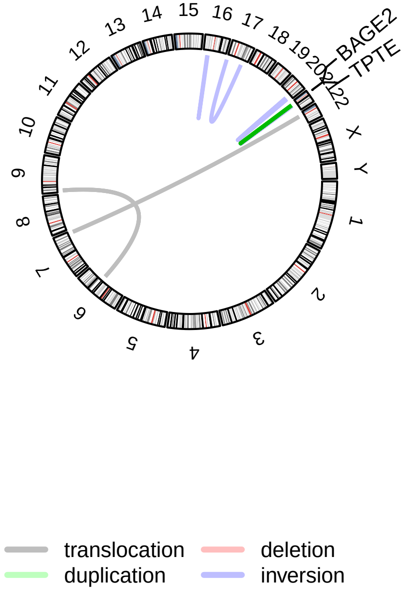




SUPPORTING READ COUNT

Split reads = 4
Discordant mates = 0

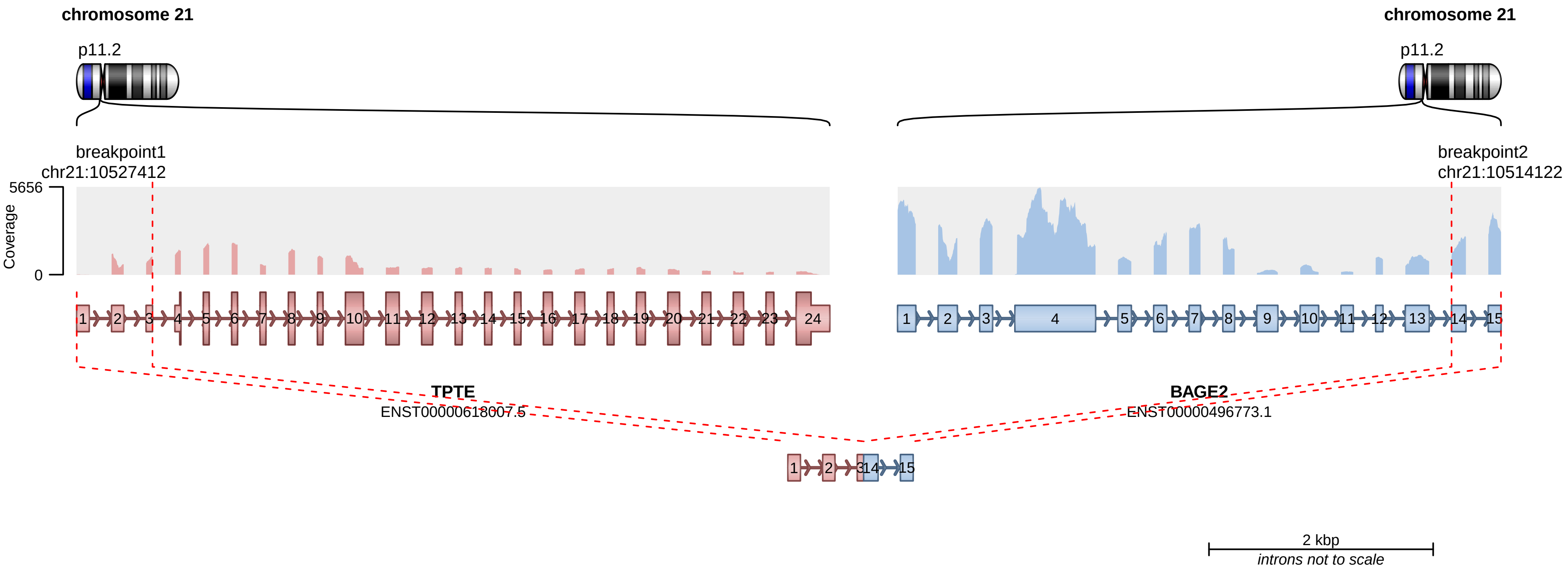


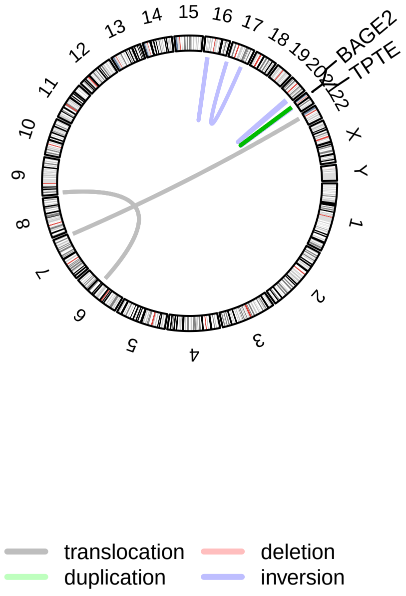


No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

Split reads = 17
Discordant mates = 0

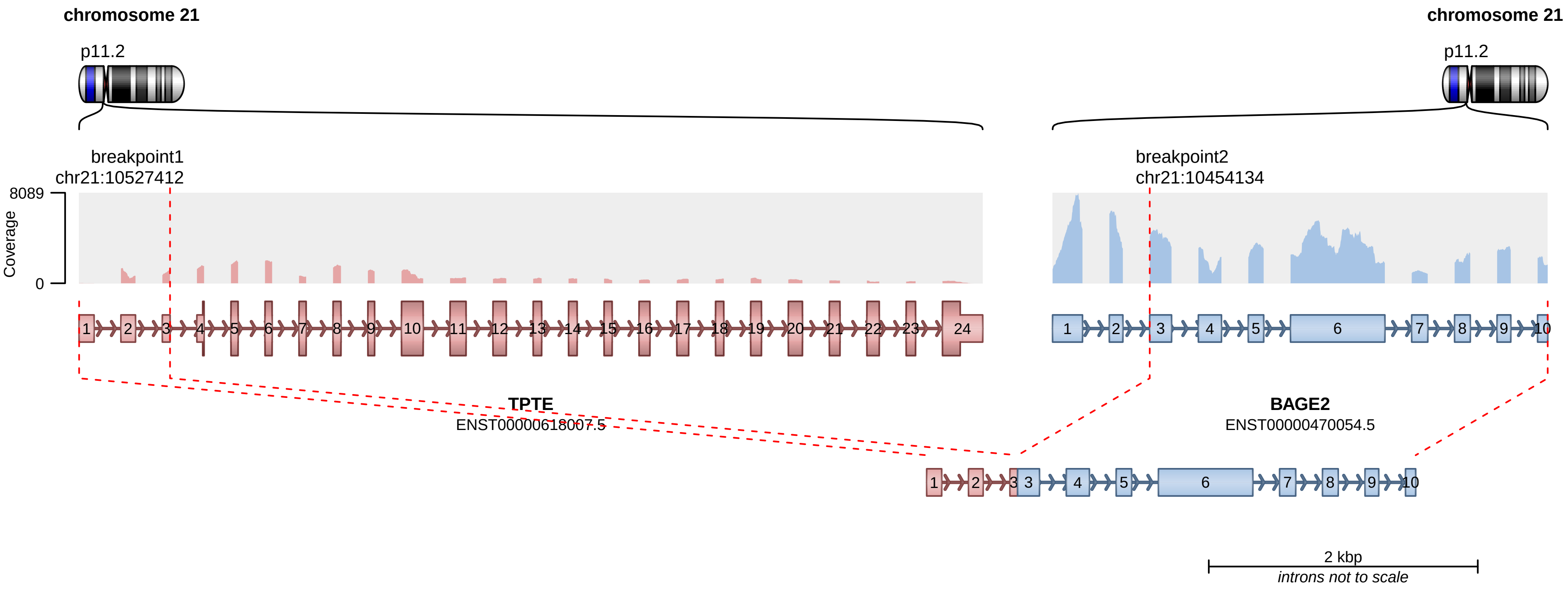


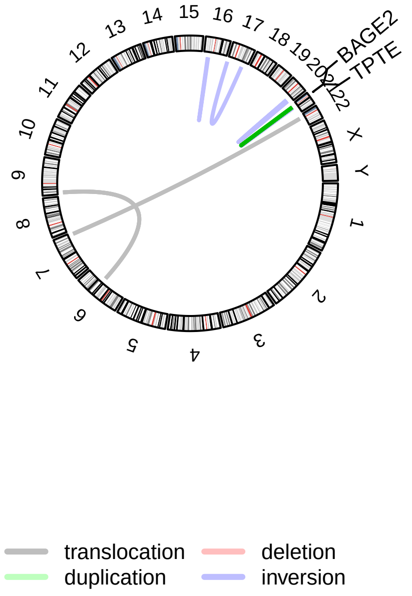


No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

Split reads = 17
Discordant mates = 0

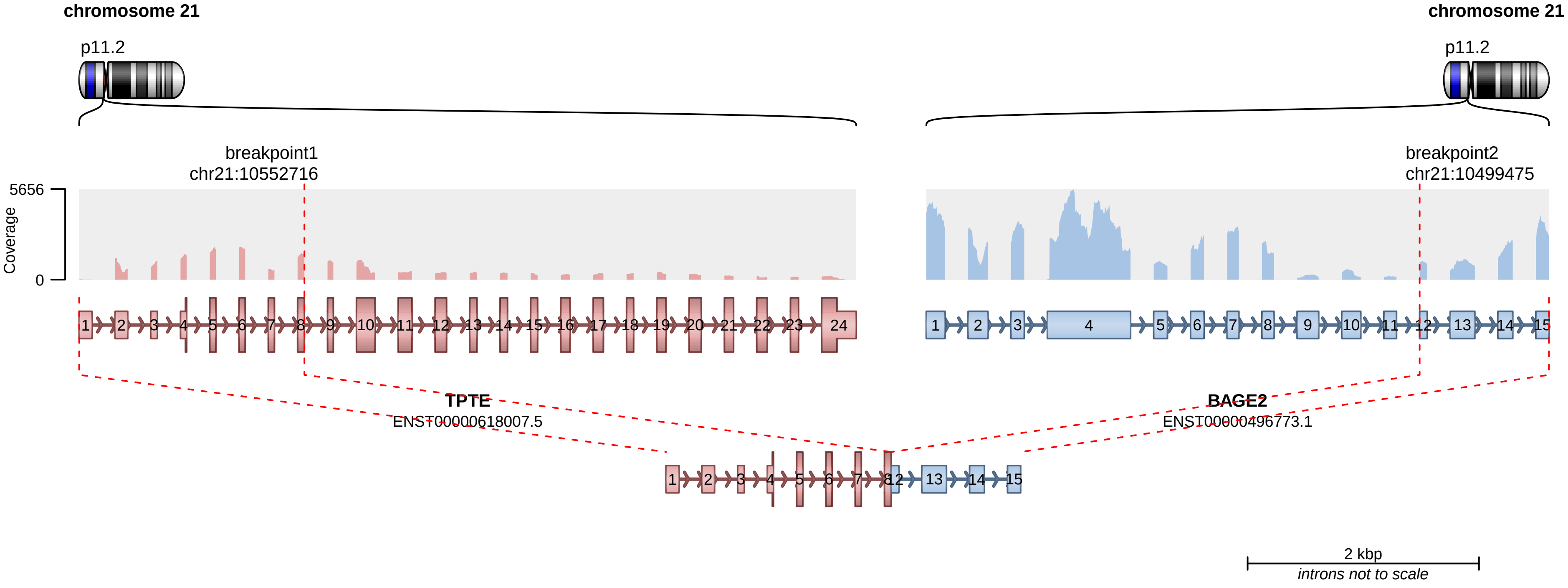


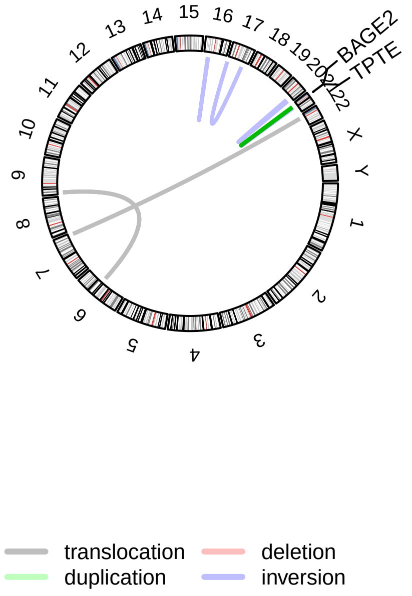


No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

Split reads = 9
Discordant mates = 0

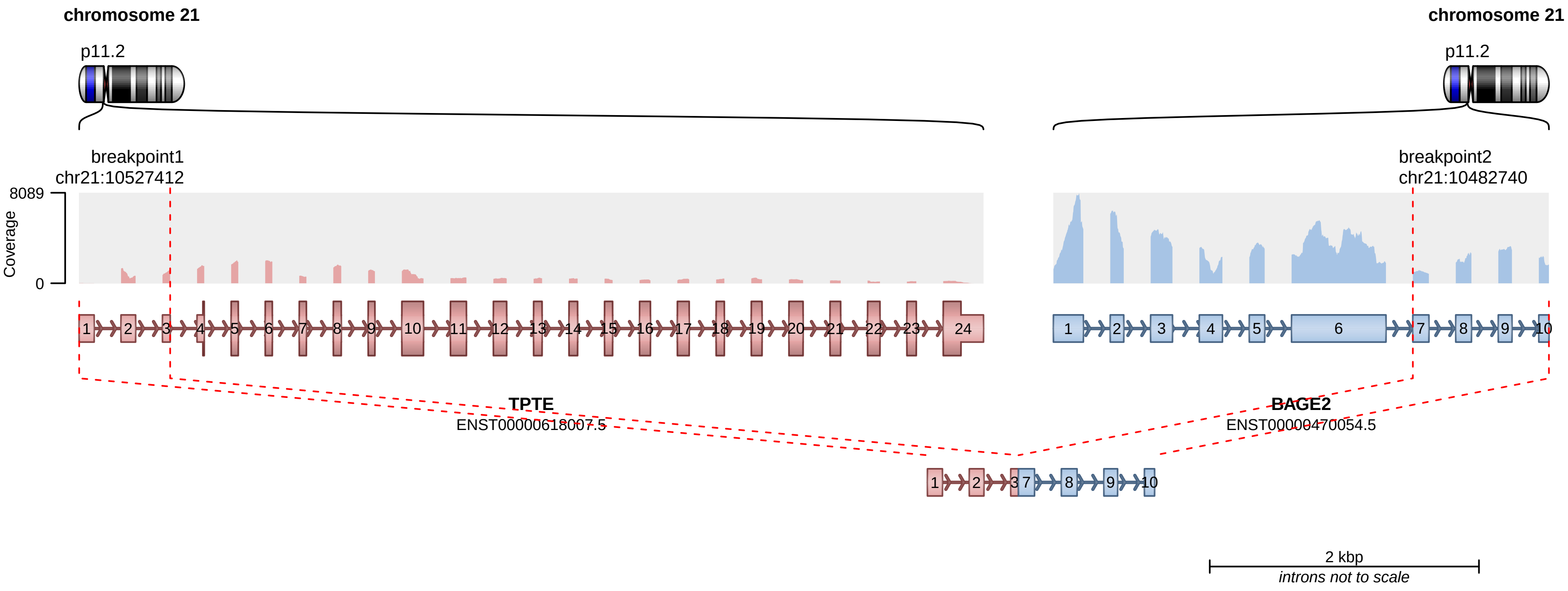


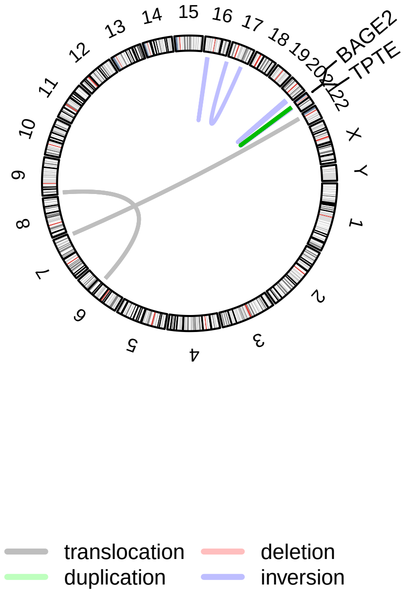


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 6
Discordant mates = 0

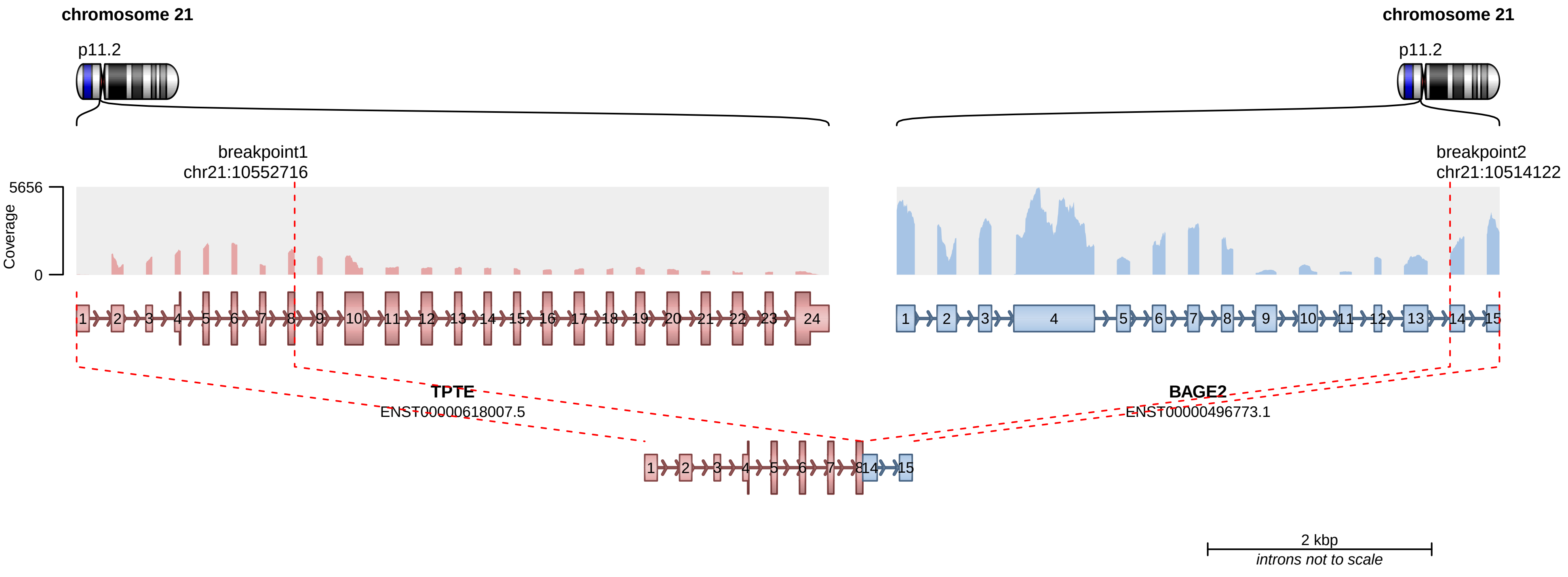


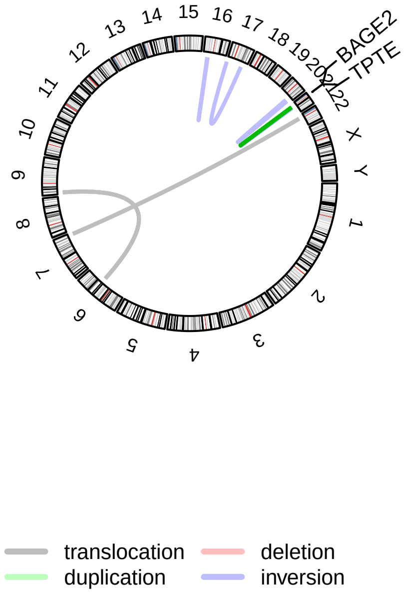


No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 0

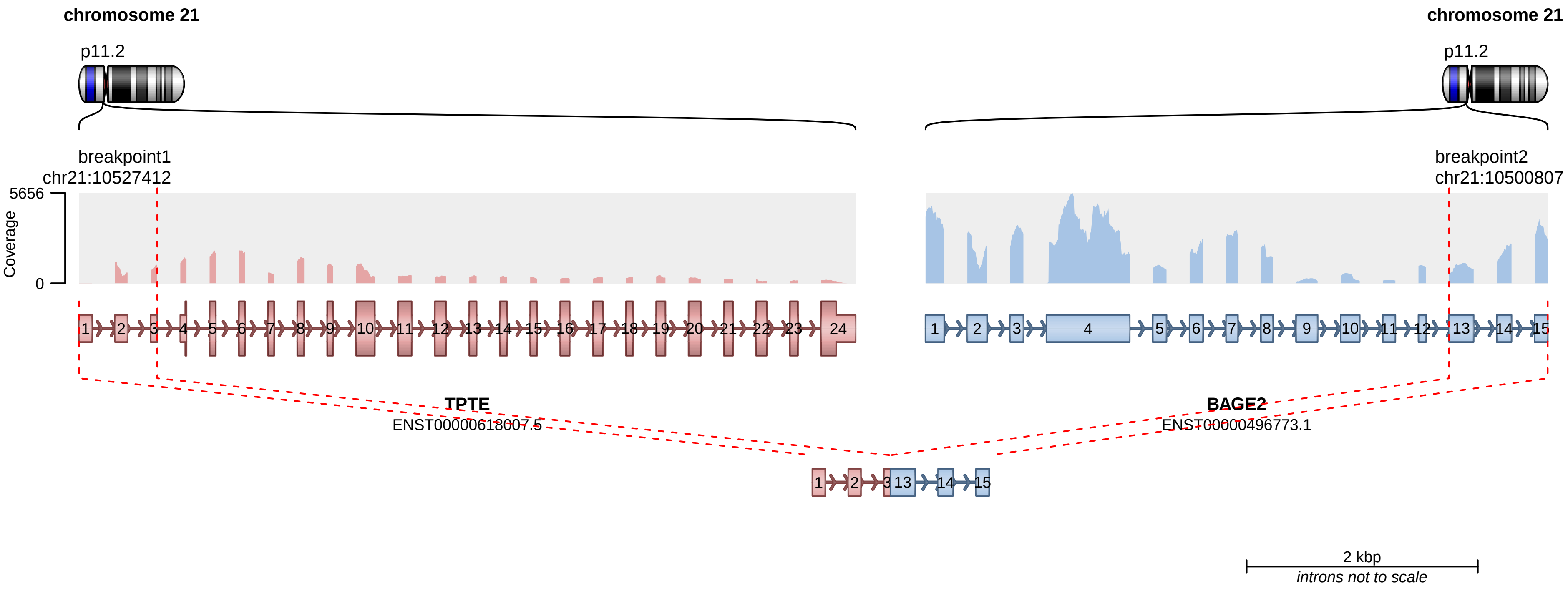


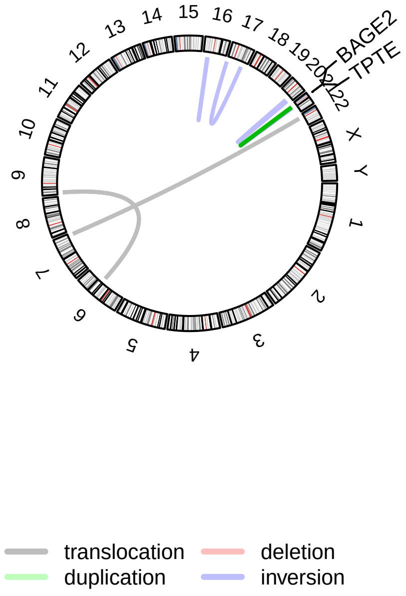


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

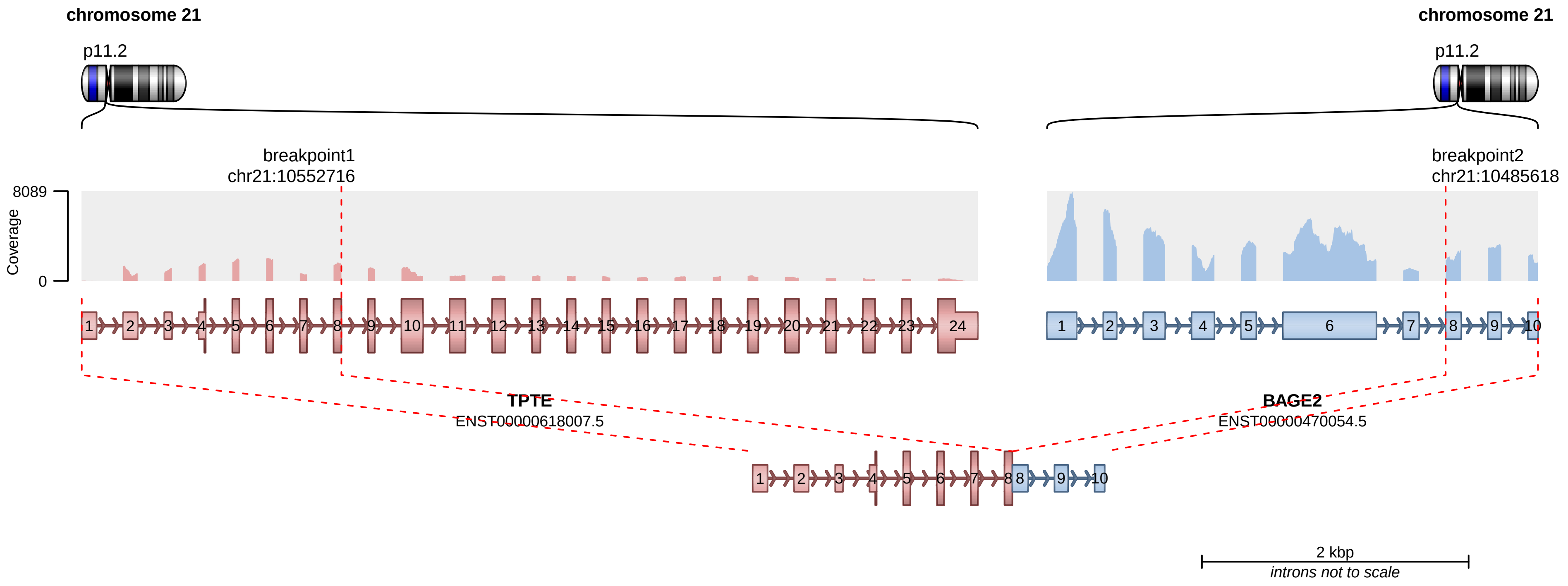


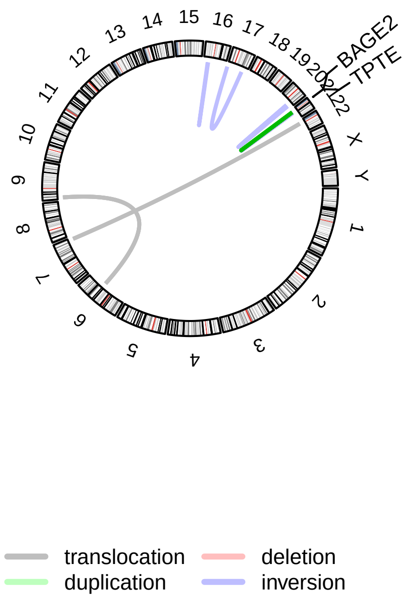


No coding regions retained in fusion transcript.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

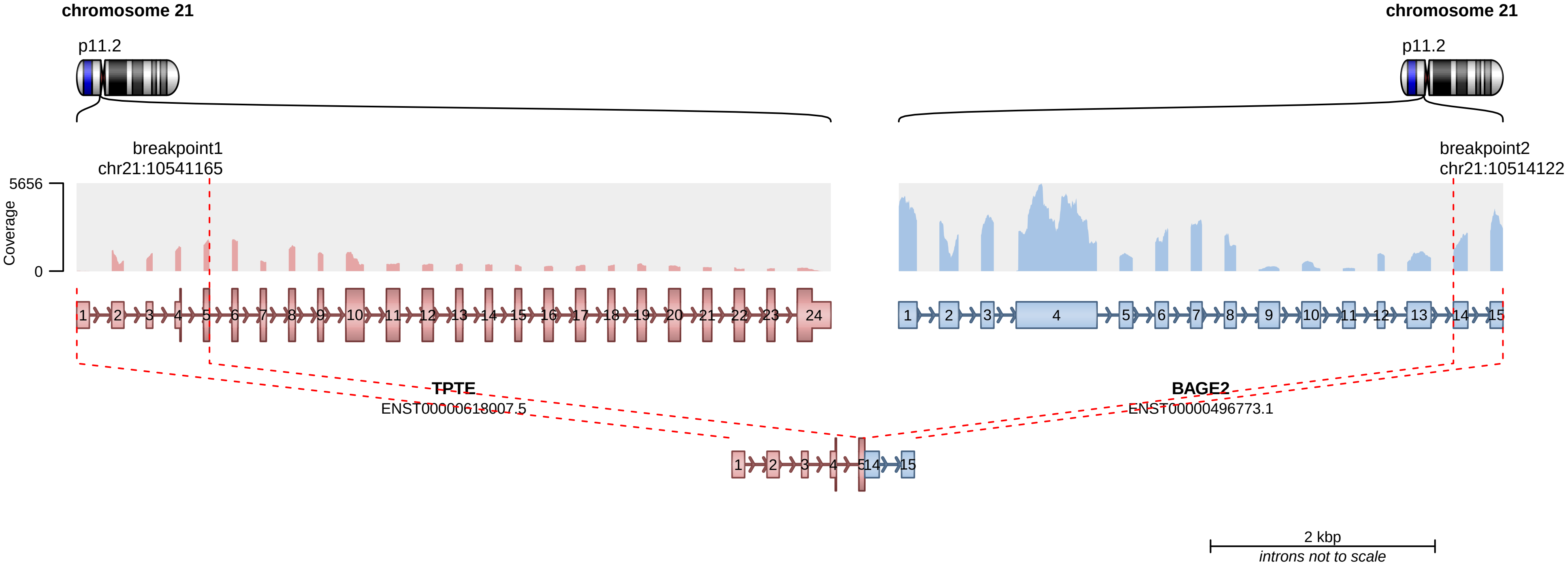


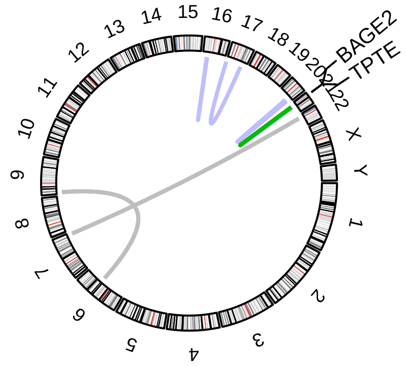


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0



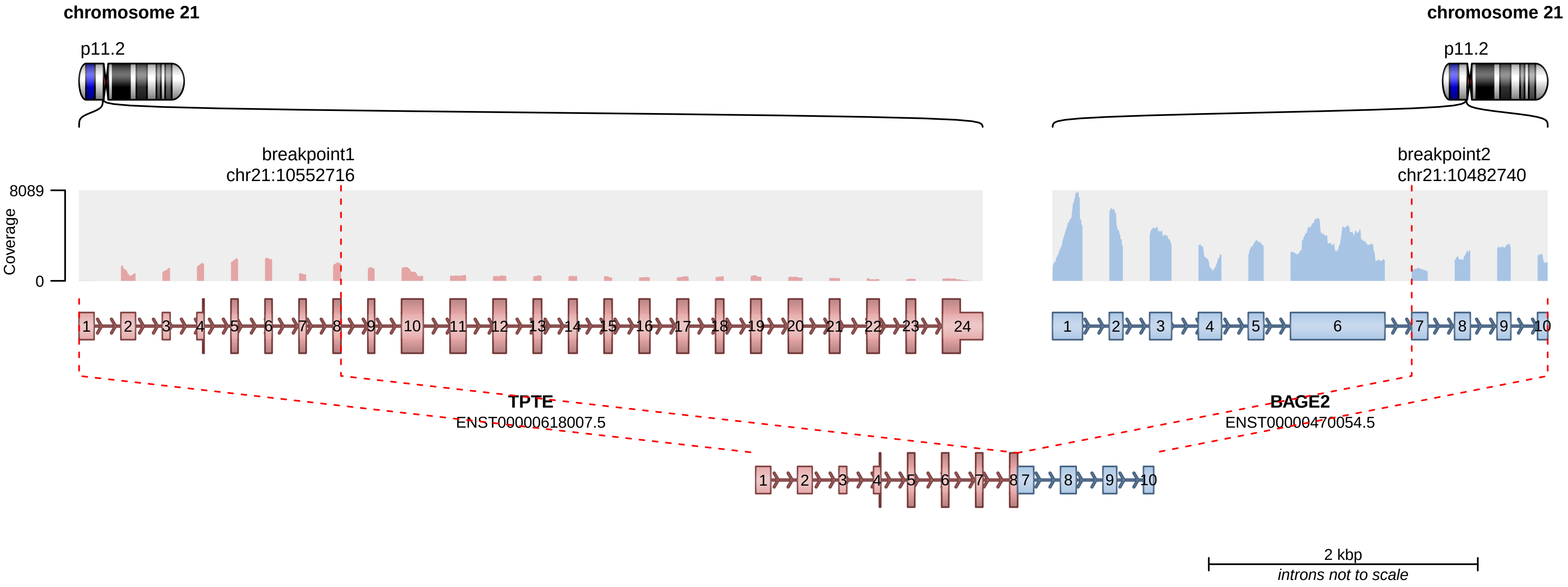


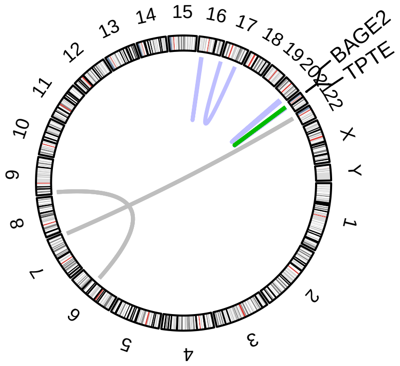
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 0

— translocation — deletion
— duplication — inversion



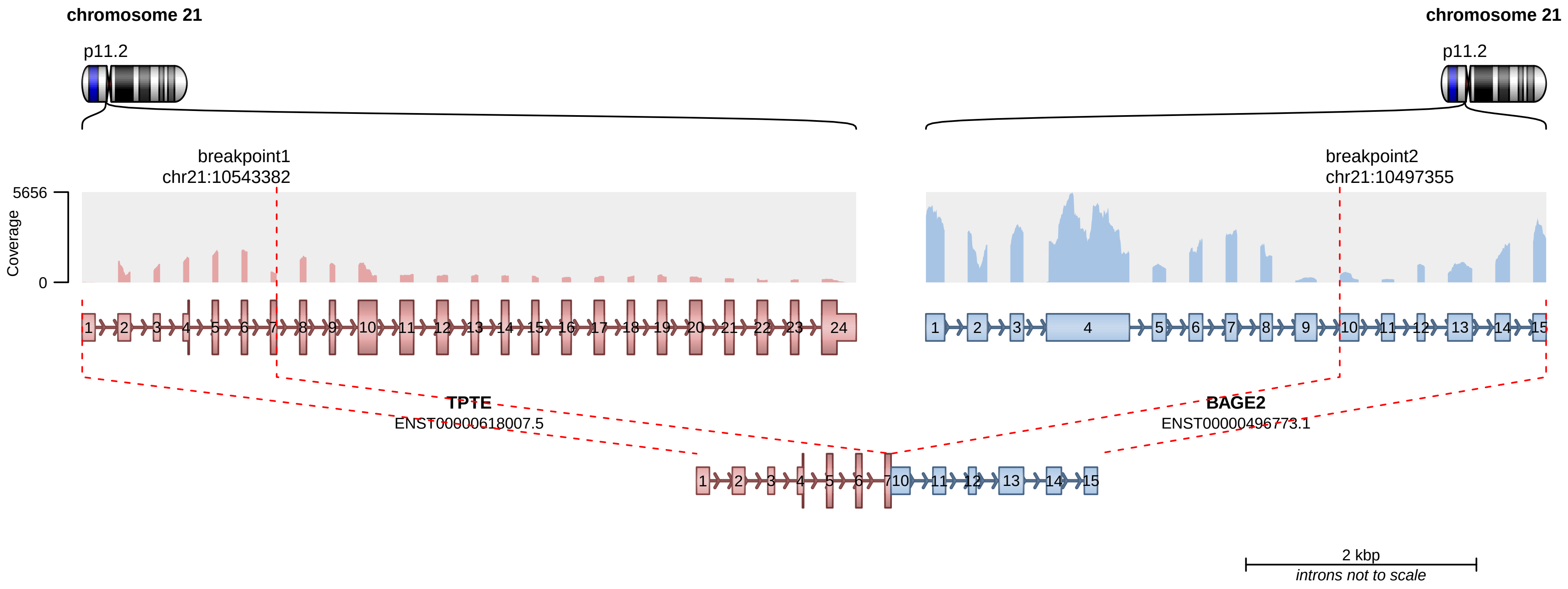


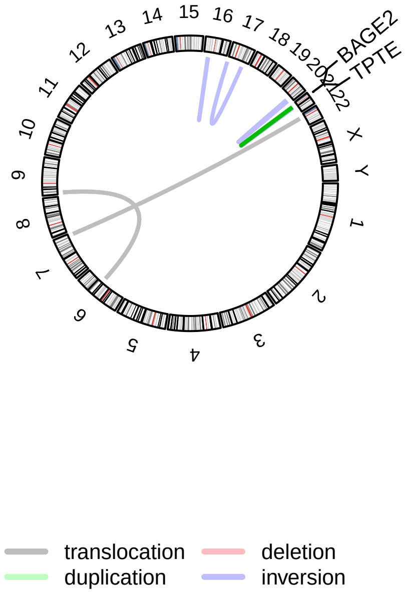
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

▬ translocation ▬ deletion
▬ duplication ▬ inversion

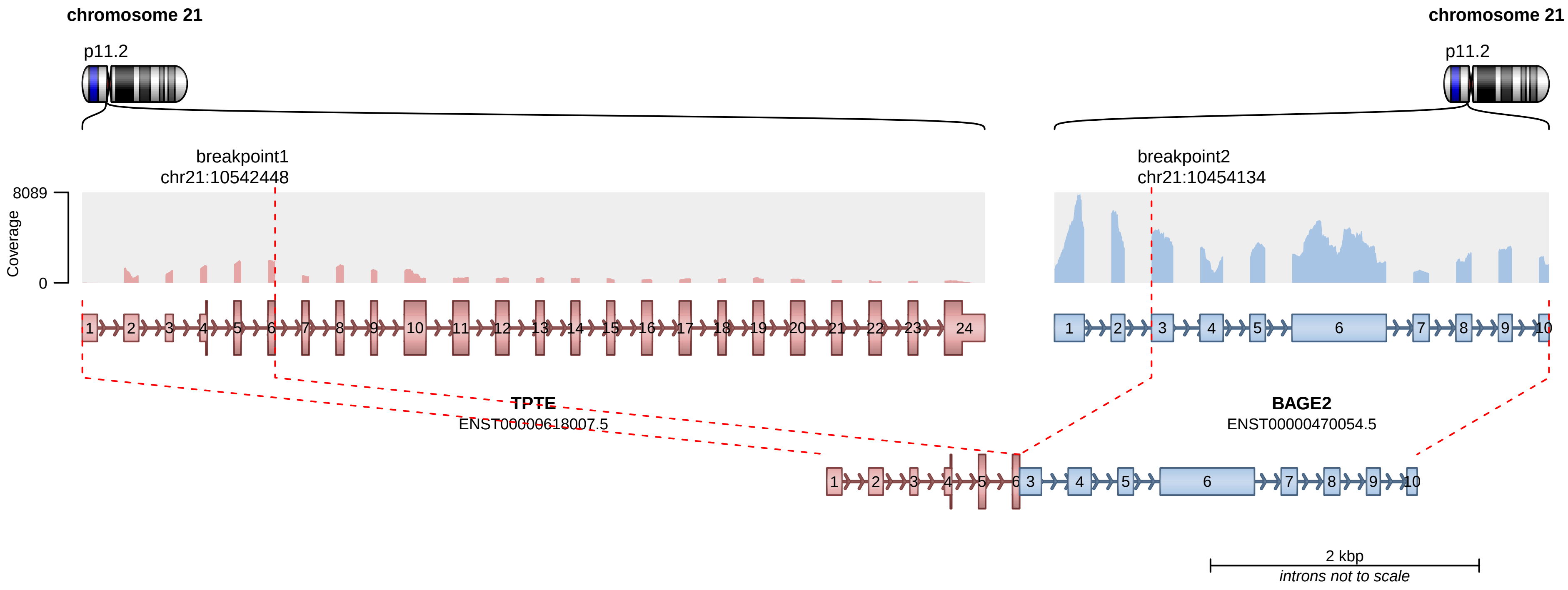


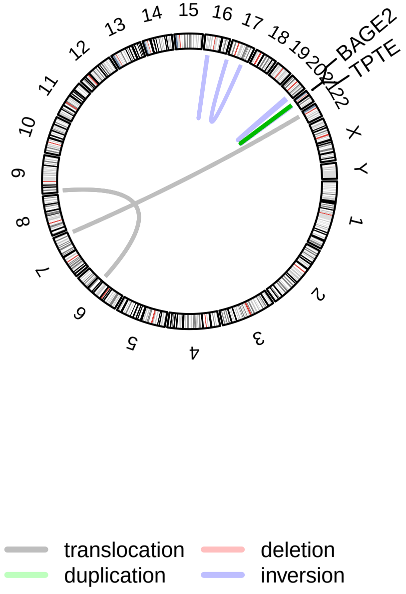


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

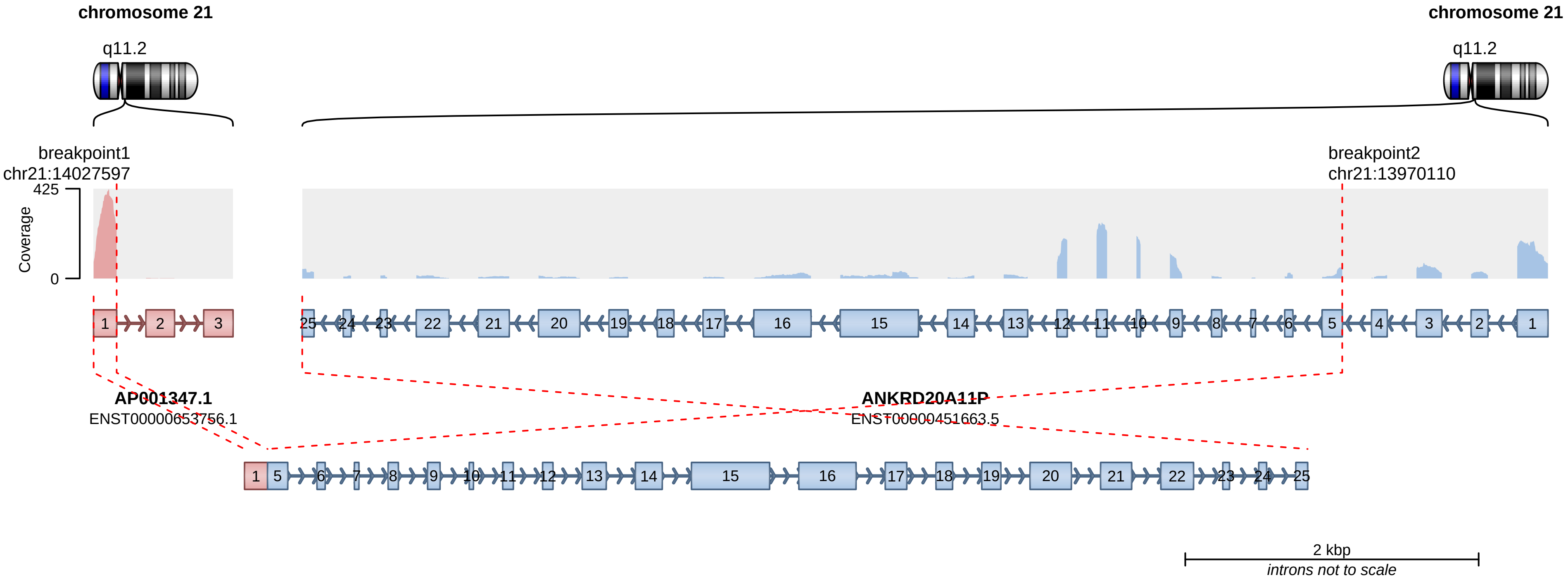


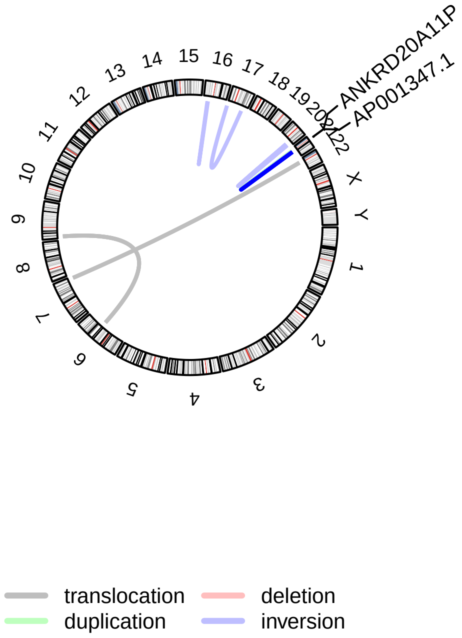


No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads = 1
Discordant mates = 0

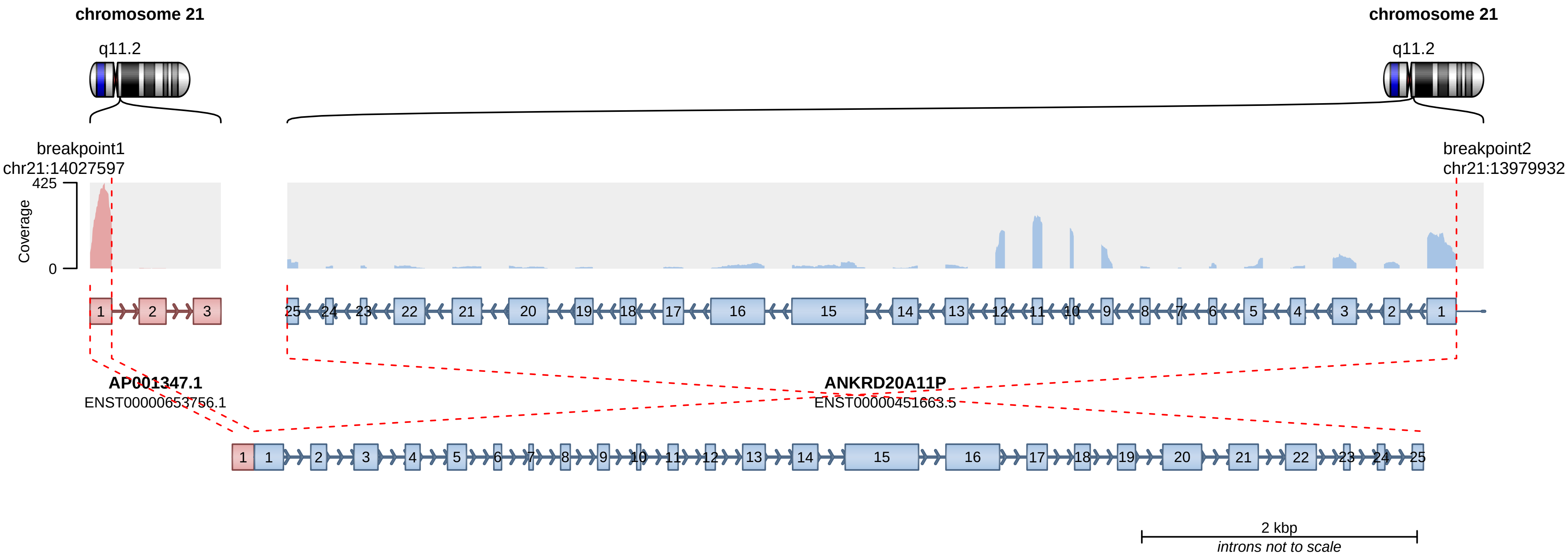


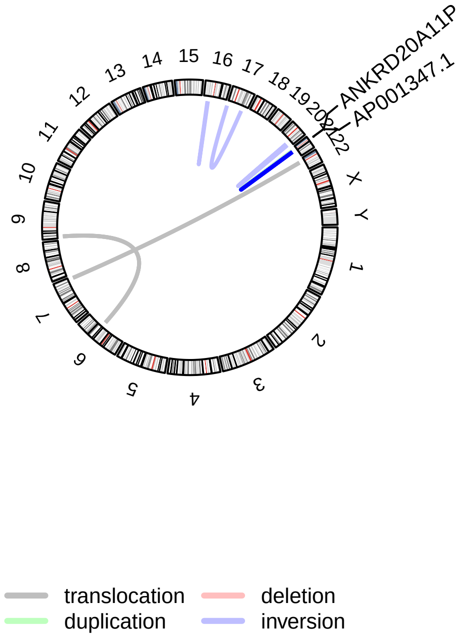


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 7
Discordant mates = 1

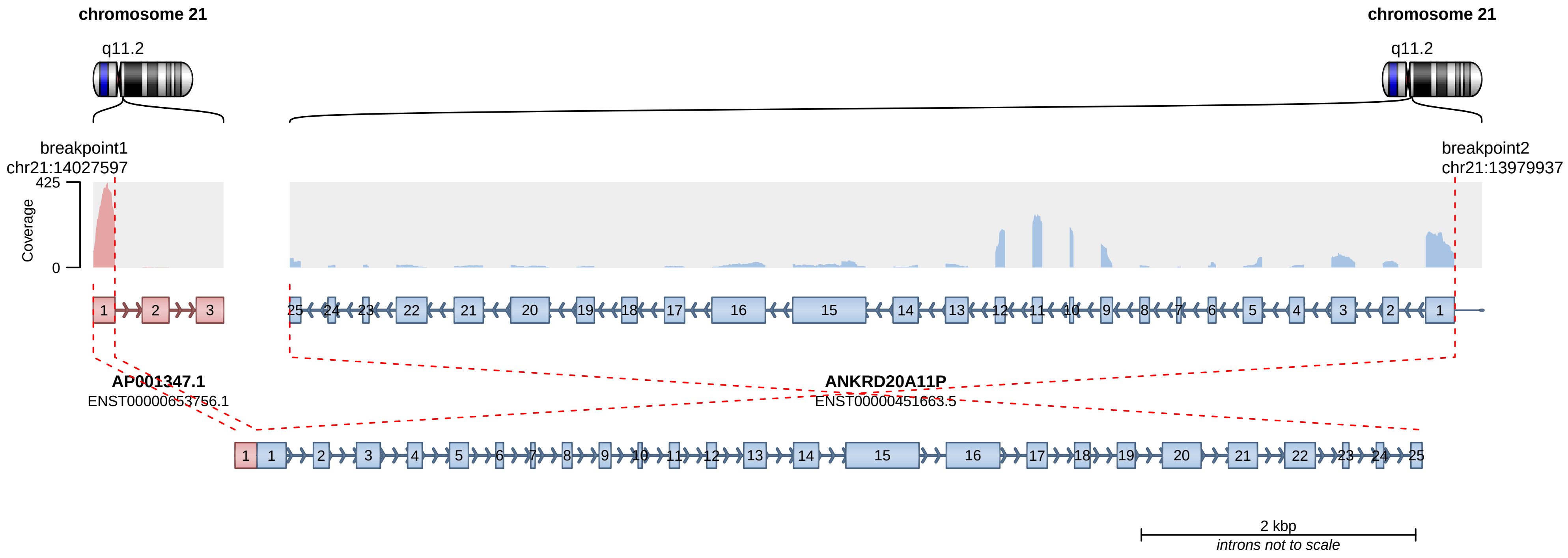


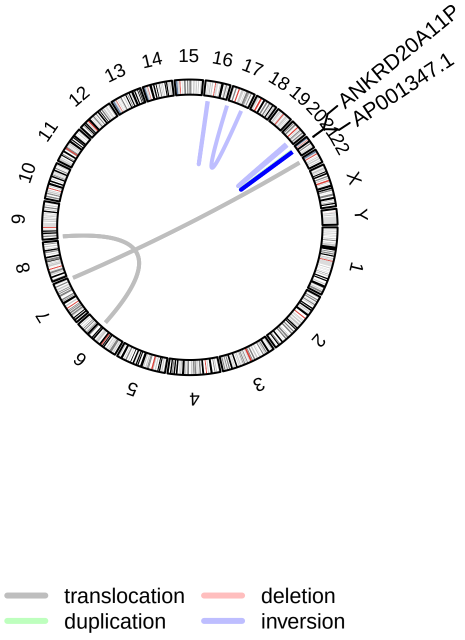


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 6
Discordant mates = 2

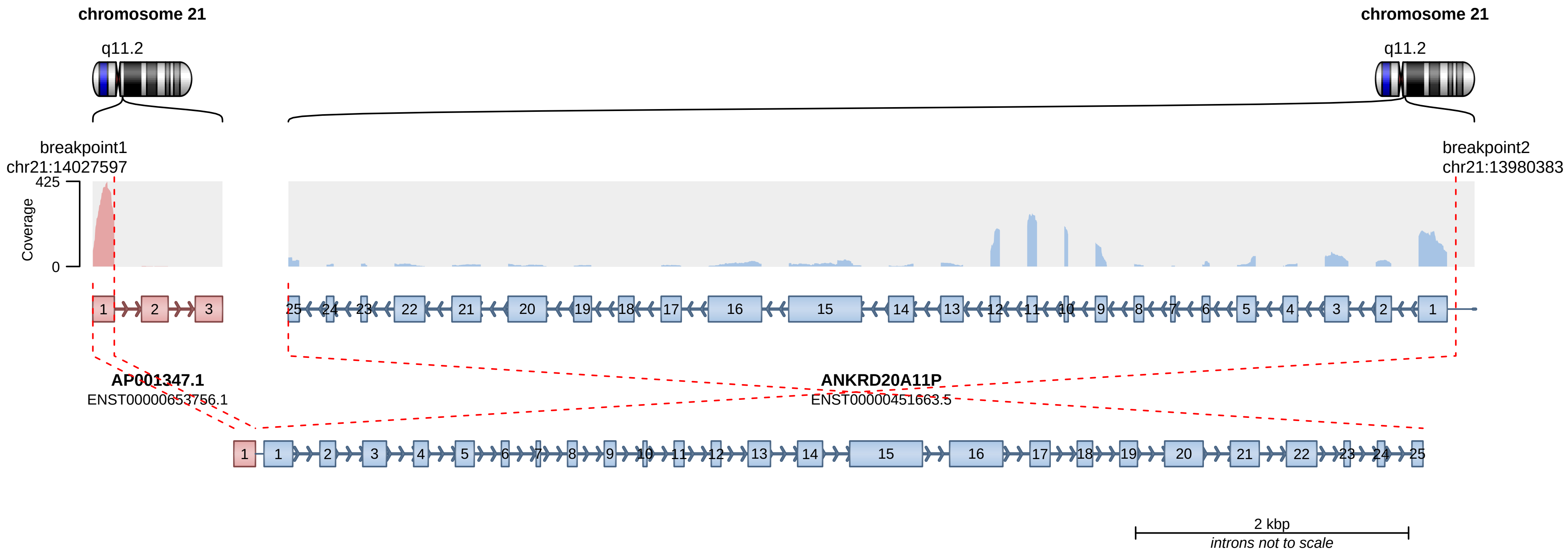


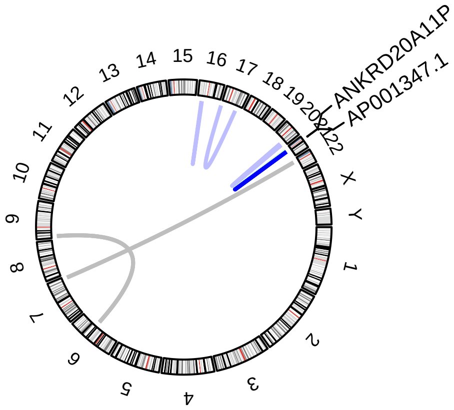


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 2



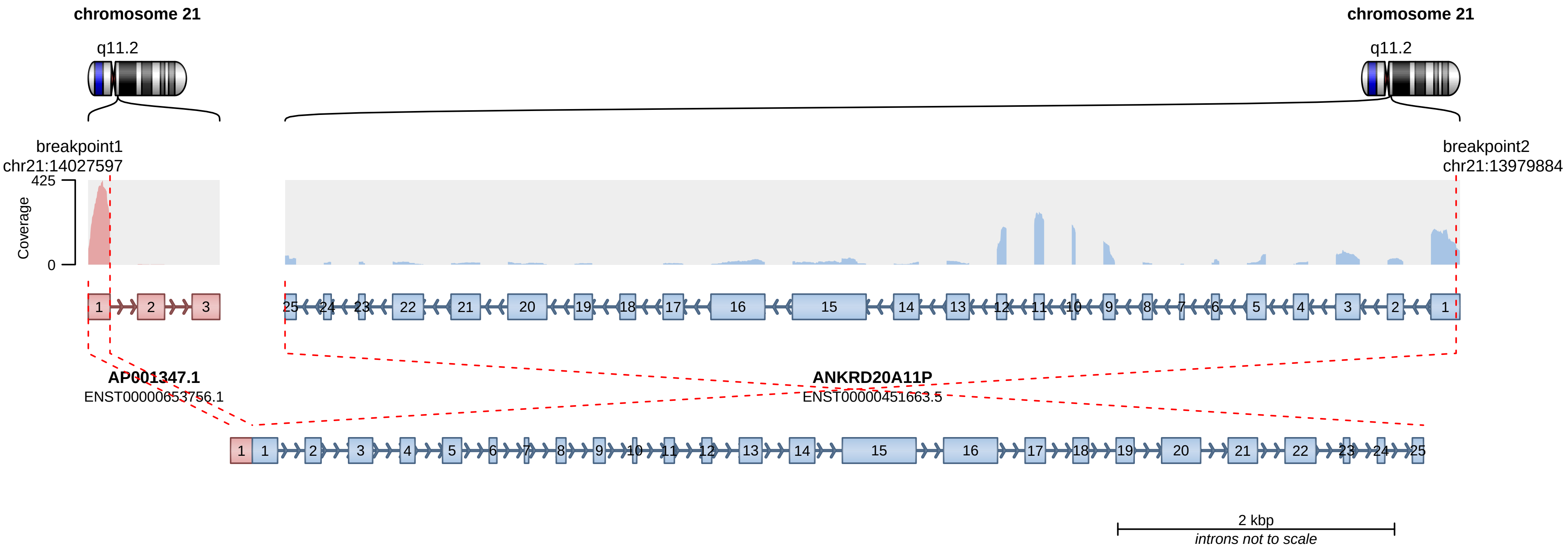


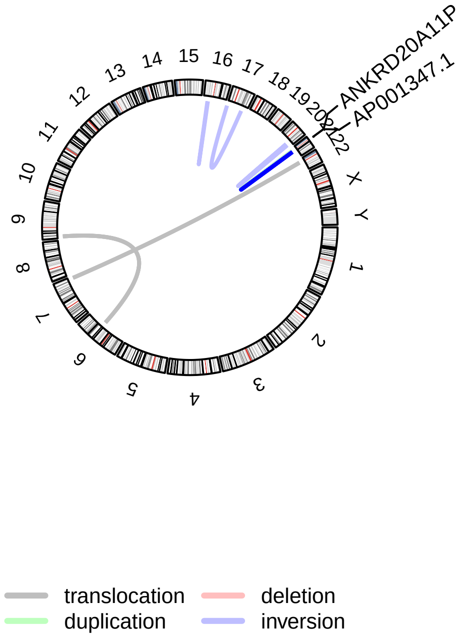
Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 2

▬ translocation ▬ deletion
▬ duplication ▬ inversion

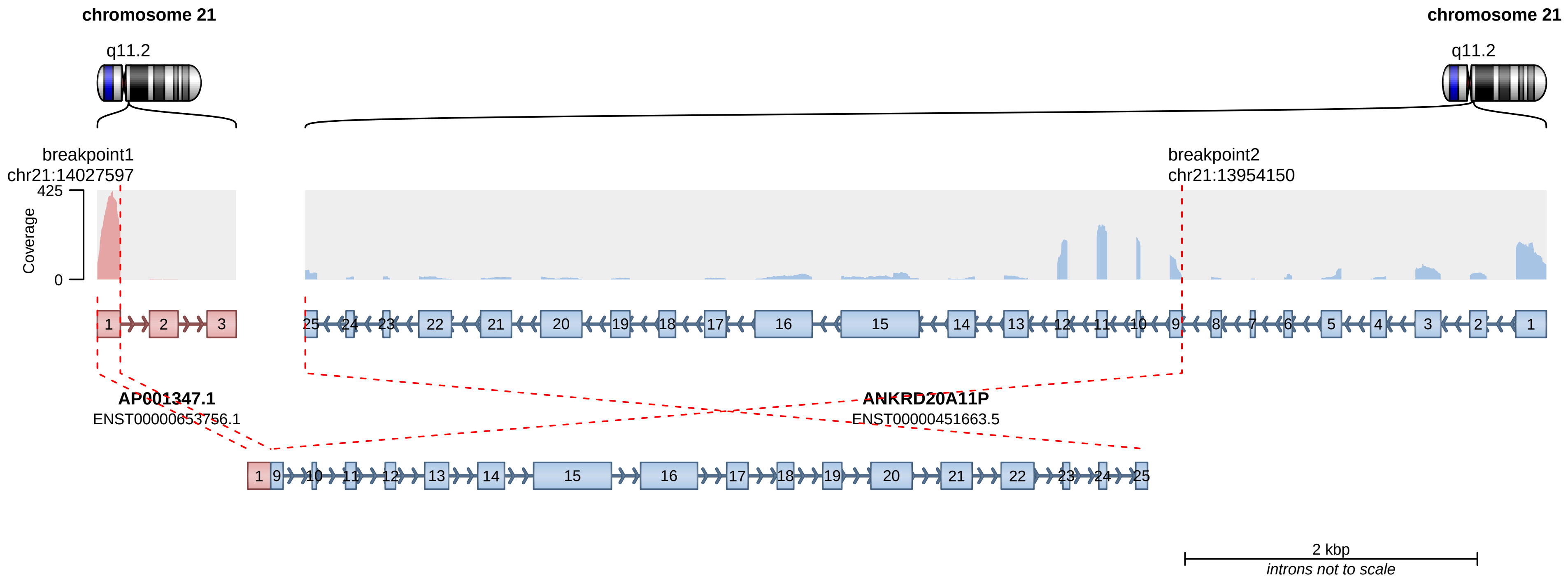


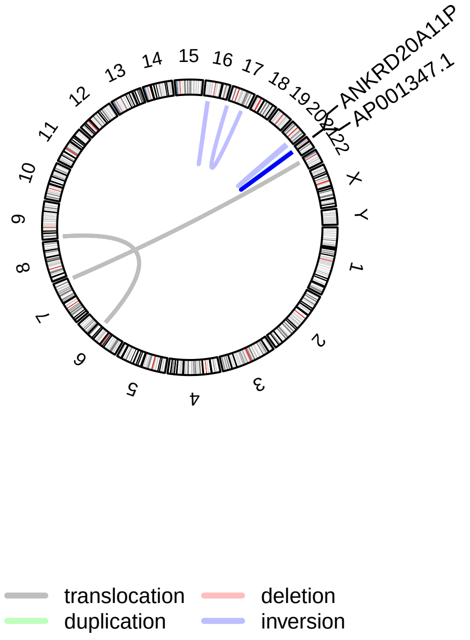


Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 3
Discordant mates = 1





Genes are not protein−coding.

SUPPORTING READ COUNT

Split reads = 2
Discordant mates = 1