

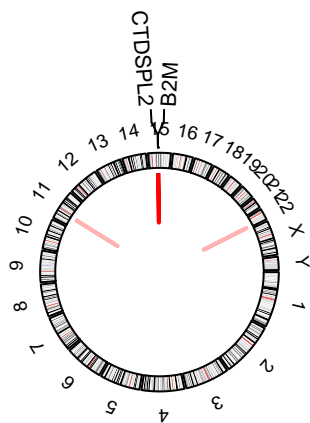
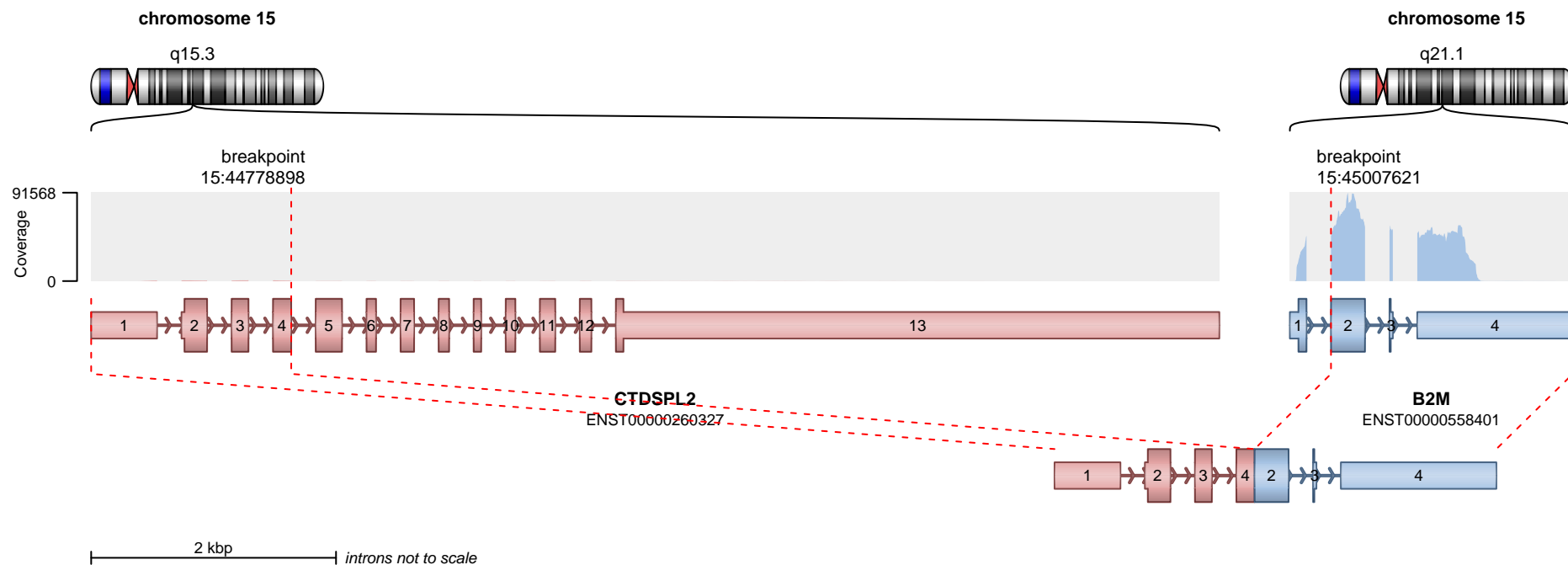
No protein domains retained in fusion.

SUPPORTING READ COUNT

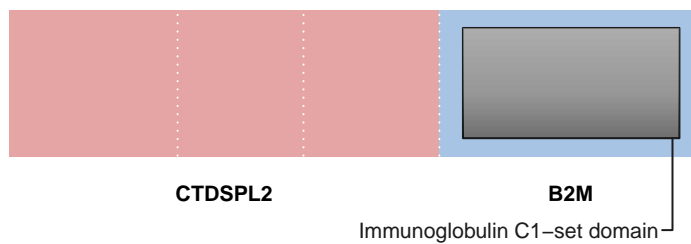
Split reads in TTC38 = 4

Split reads in ATXN10(10077),WI2-85898F10.1(20615) = 7

Discordant mates = 3



RETAINED PROTEIN DOMAINS reading frame unclear

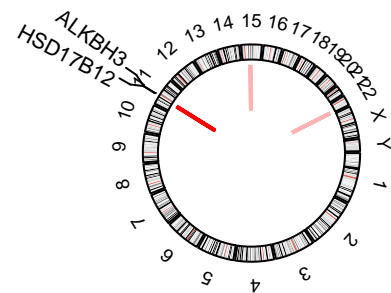
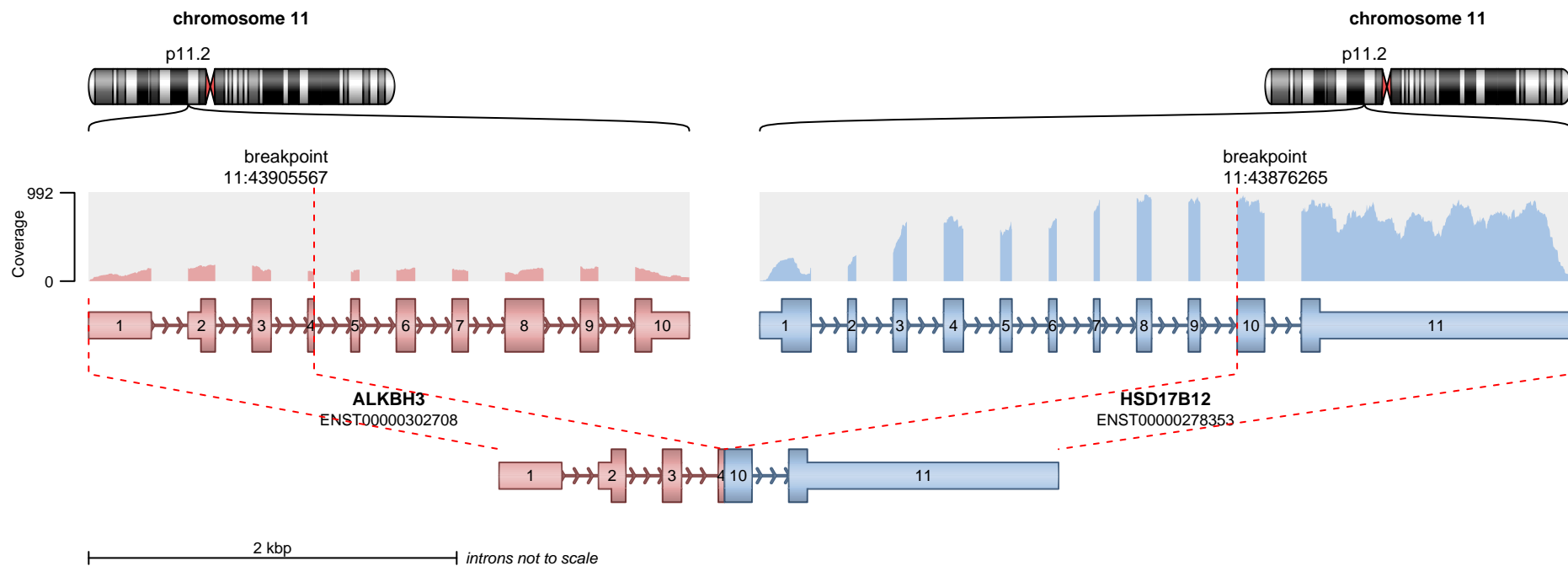


SUPPORTING READ COUNT

Split reads in CTDSPL2 = 36

Split reads in B2M = 65

Discordant mates = 13



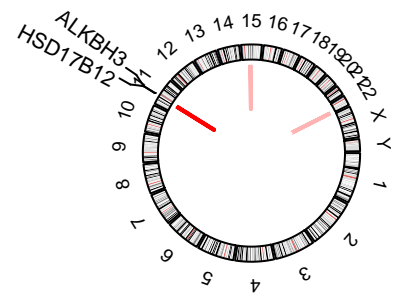
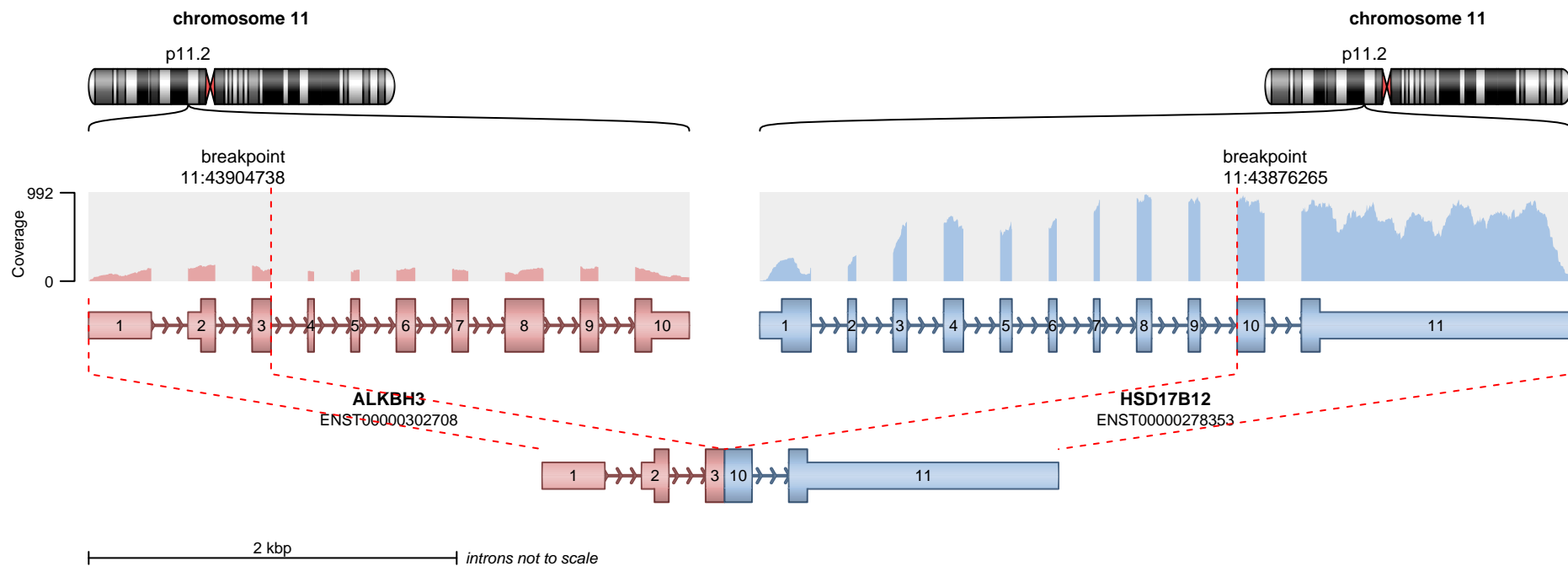
No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads in ALKBH3 = 1

Split reads in HSD17B12 = 1

Discordant mates = 0



No protein domains retained in fusion.

SUPPORTING READ COUNT

Split reads in ALKBH3 = 3

Split reads in HSD17B12 = 0

Discordant mates = 0