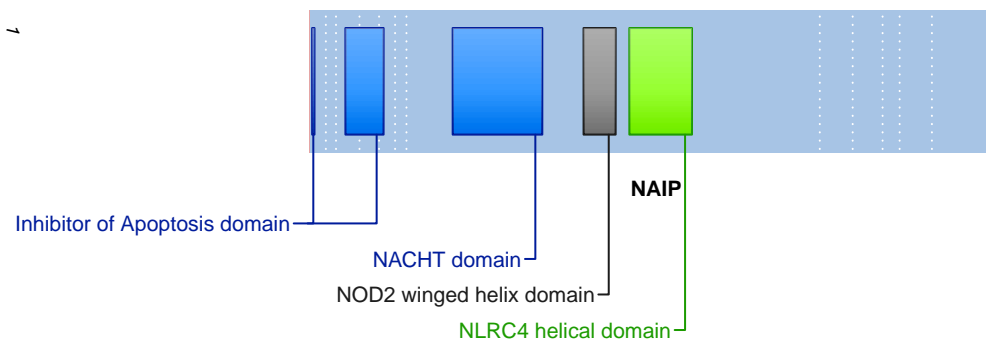


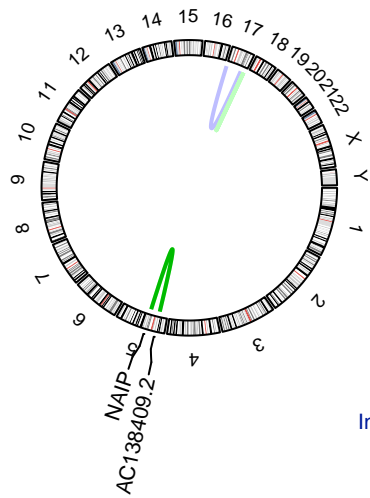
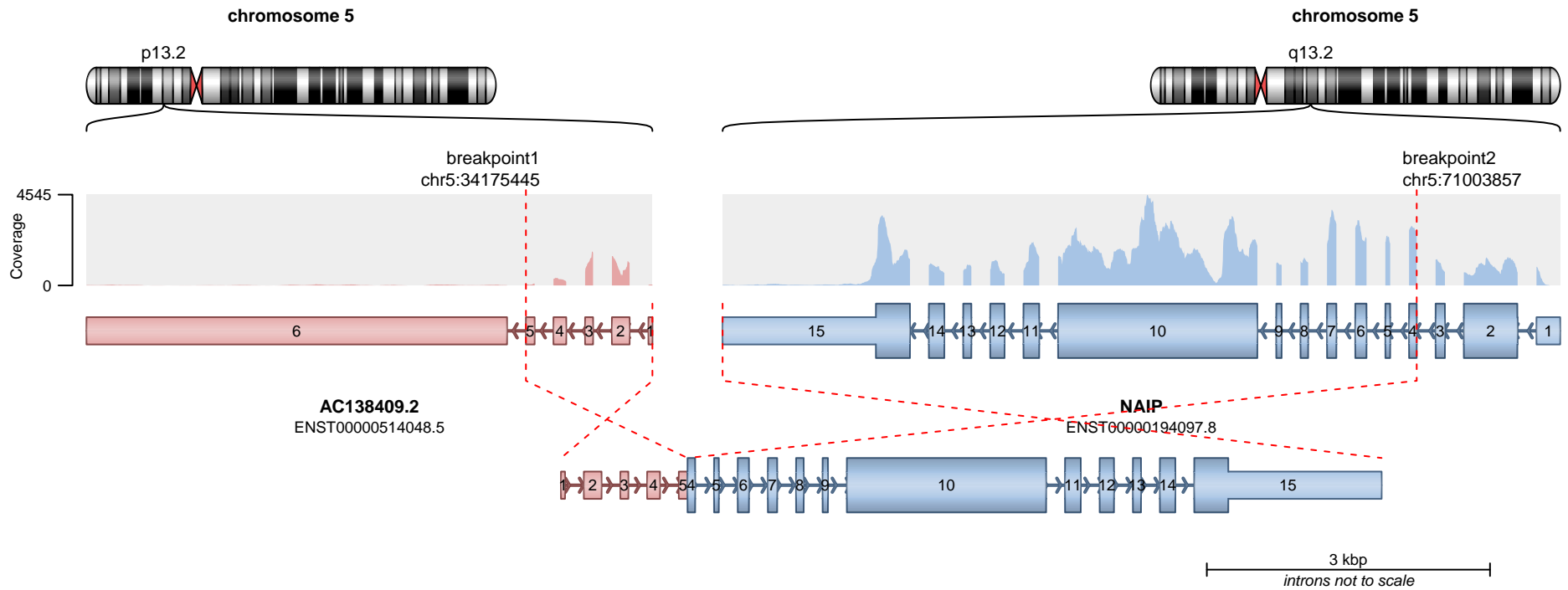
— translocation — deletion  
— duplication — inversion

**RETAINED PROTEIN DOMAINS**  
reading frame unclear



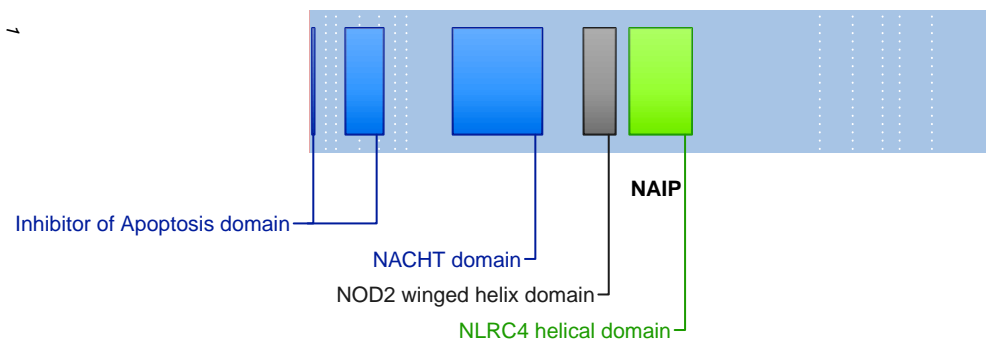
**SUPPORTING READ COUNT**

Split reads = 125  
Discordant mates = 0



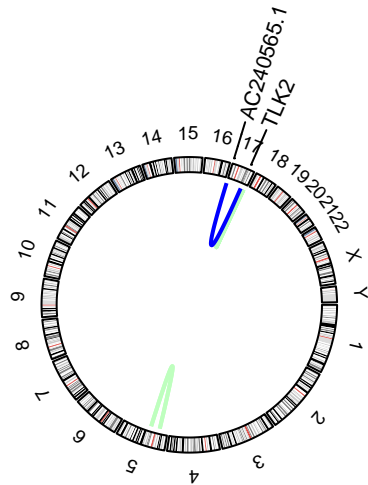
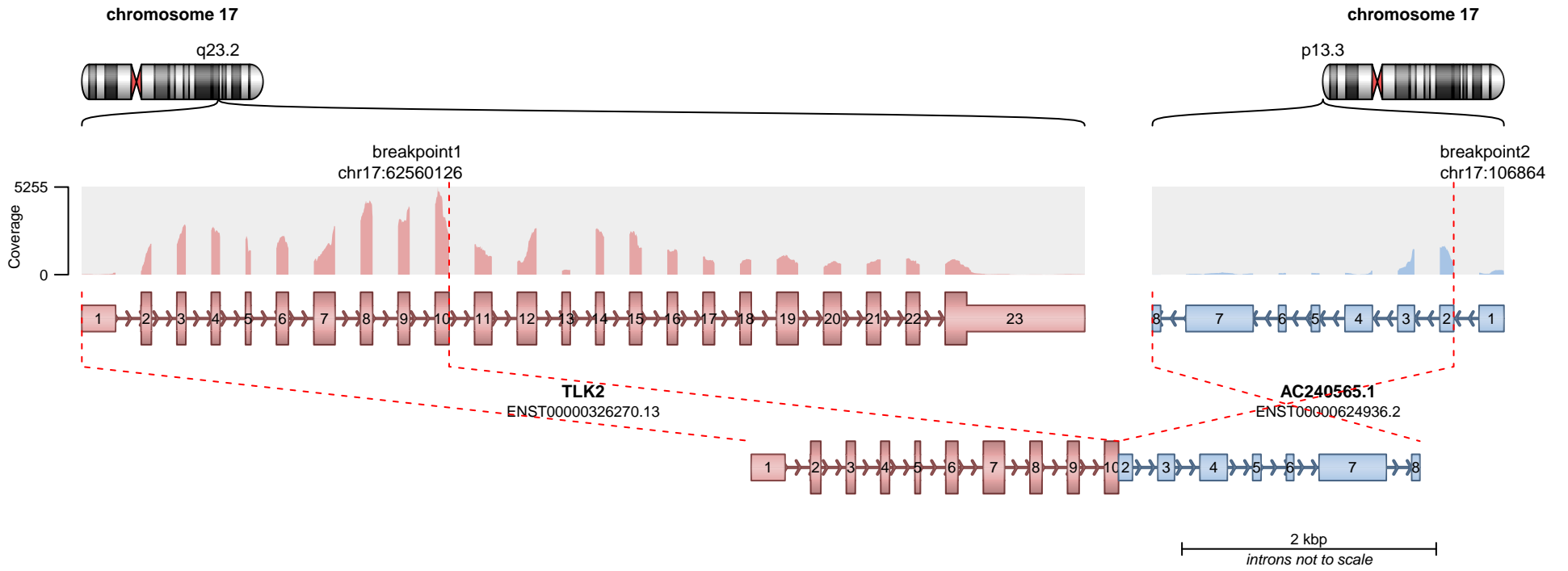
— translocation — deletion  
— duplication — inversion

**RETAINED PROTEIN DOMAINS**  
reading frame unclear



**SUPPORTING READ COUNT**

Split reads = 24  
Discordant mates = 0

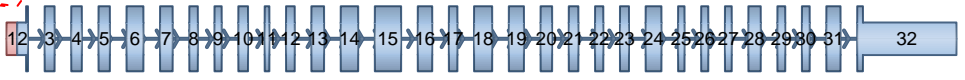
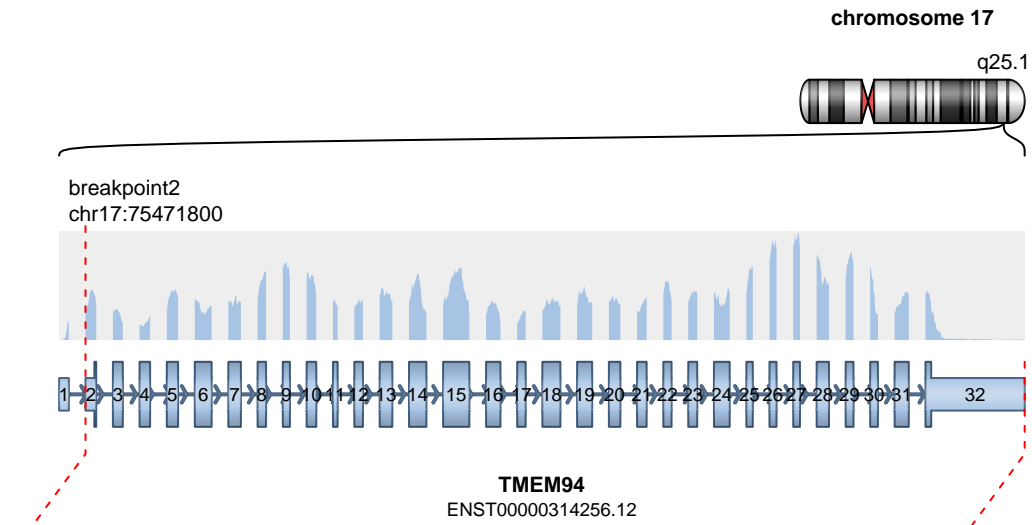
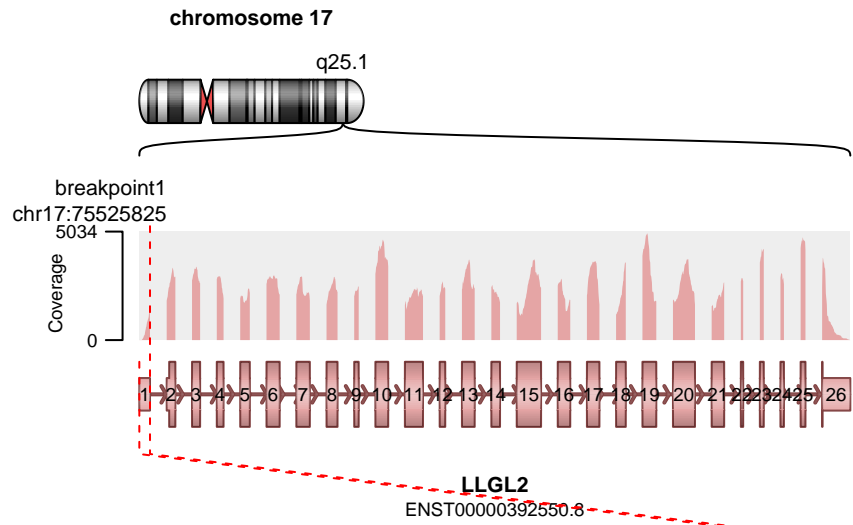


No protein domains retained in fusion.

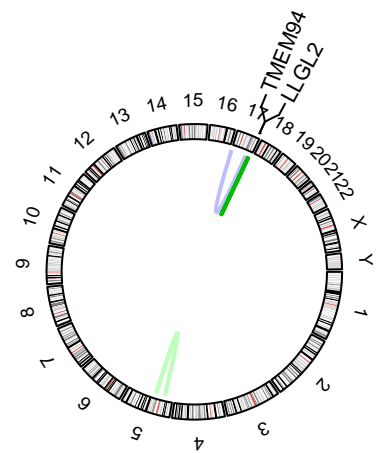
**SUPPORTING READ COUNT**

Split reads = 69  
Discordant mates = 0

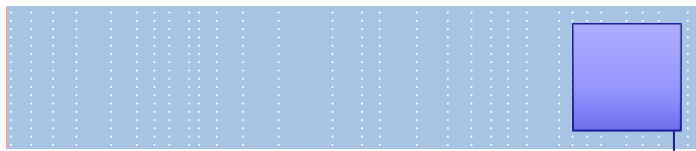
— translocation — deletion  
— duplication — inversion



4 kbp  
introns not to scale



**RETAINED PROTEIN DOMAINS**  
reading frame unclear

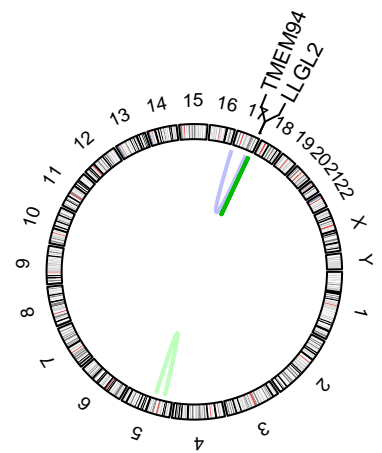
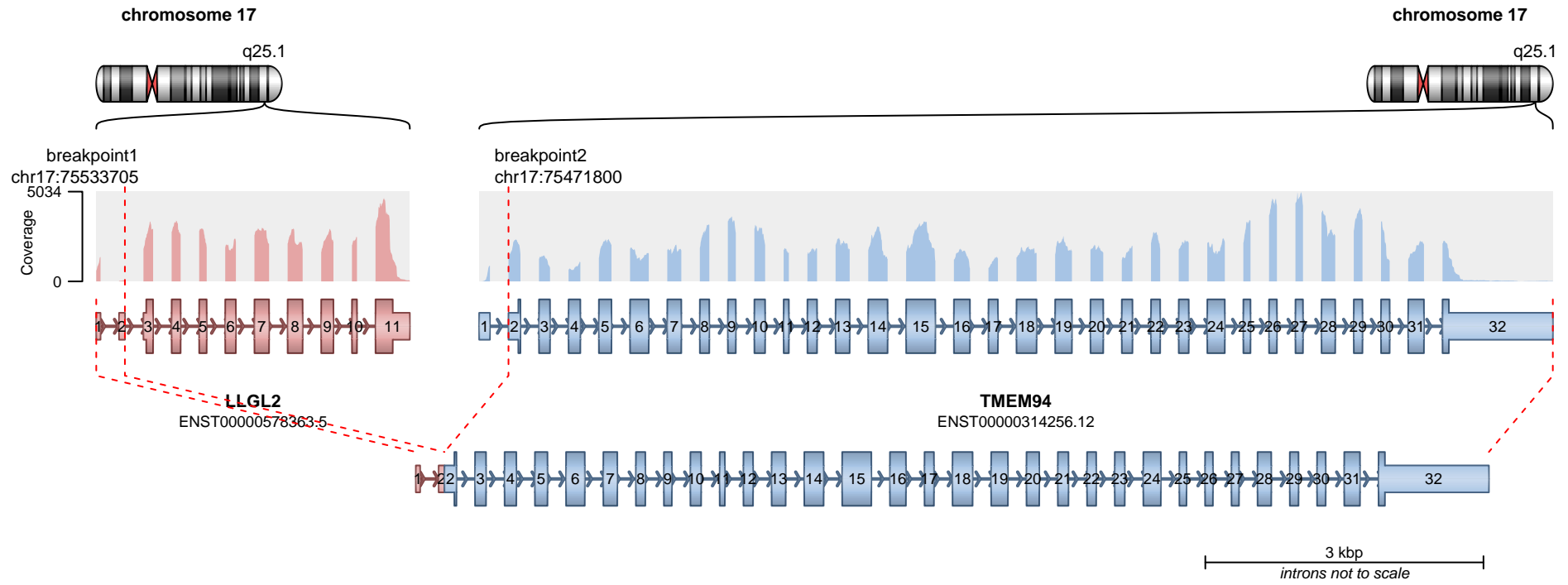


Cation transporting ATPase, C-terminus

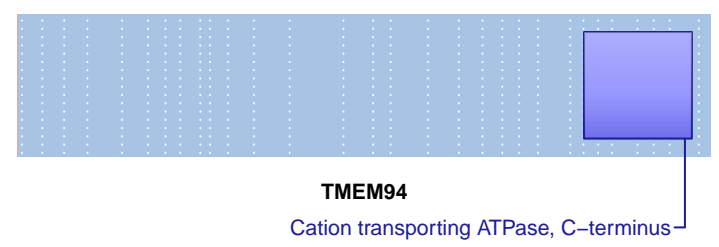
**SUPPORTING READ COUNT**

Split reads = 13  
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