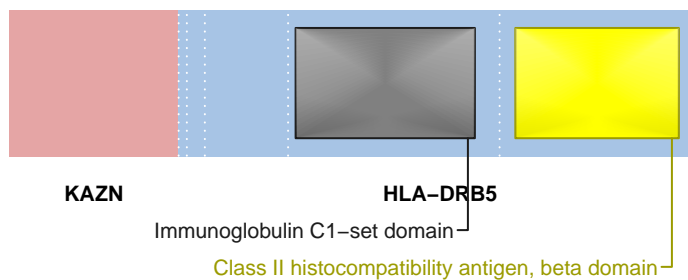
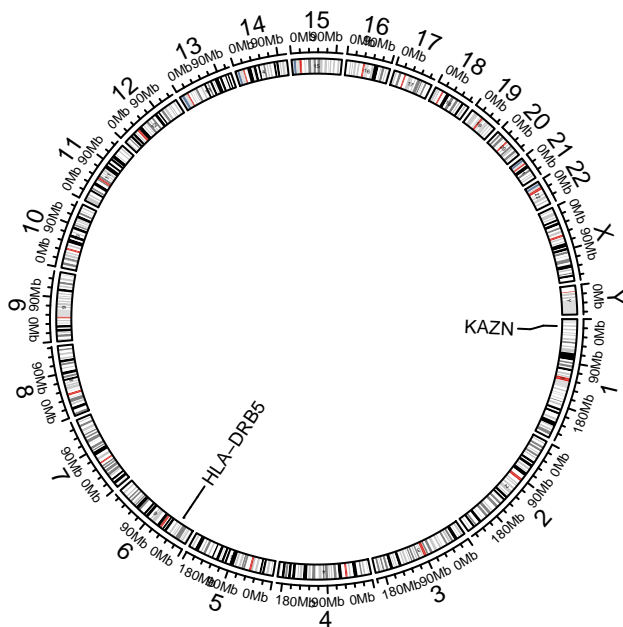


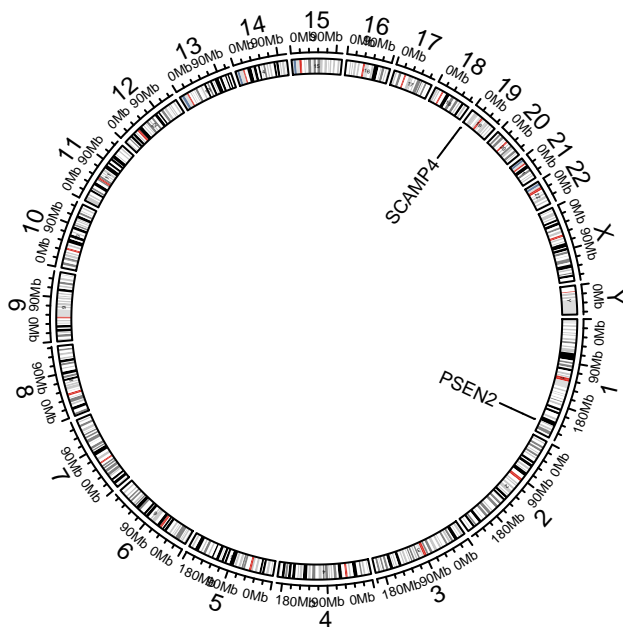
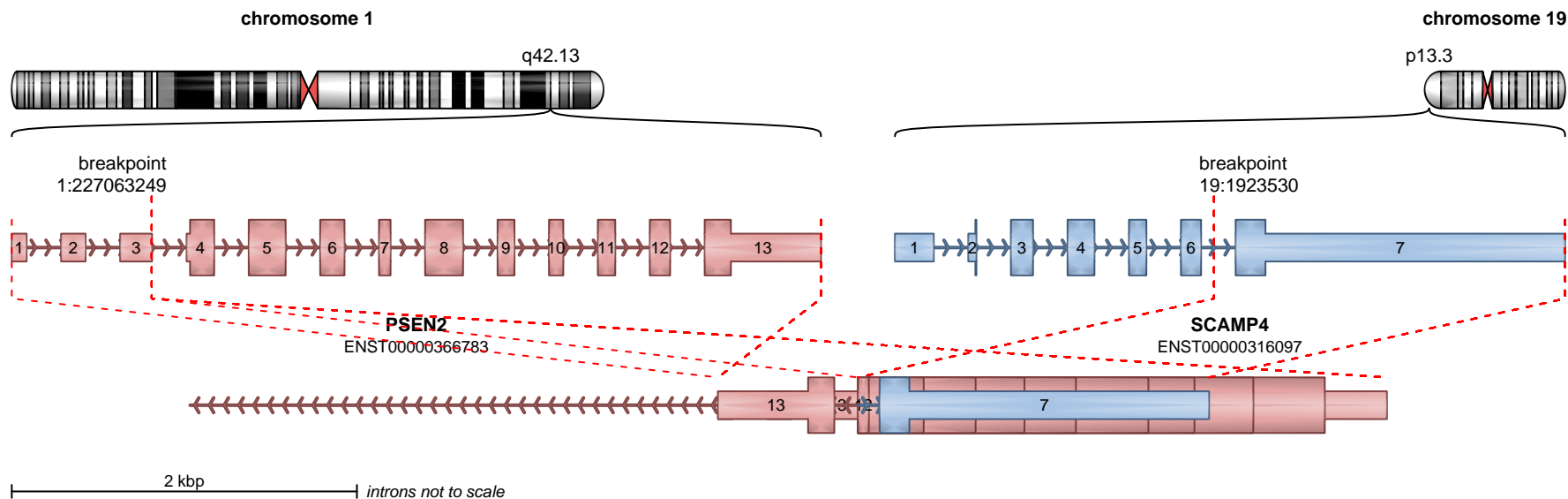
RETAINED PROTEIN DOMAINS



SUPPORTING READ COUNT

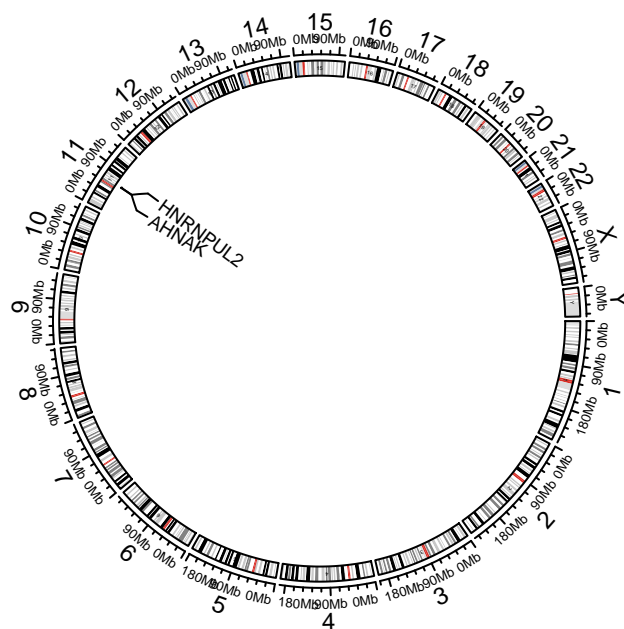
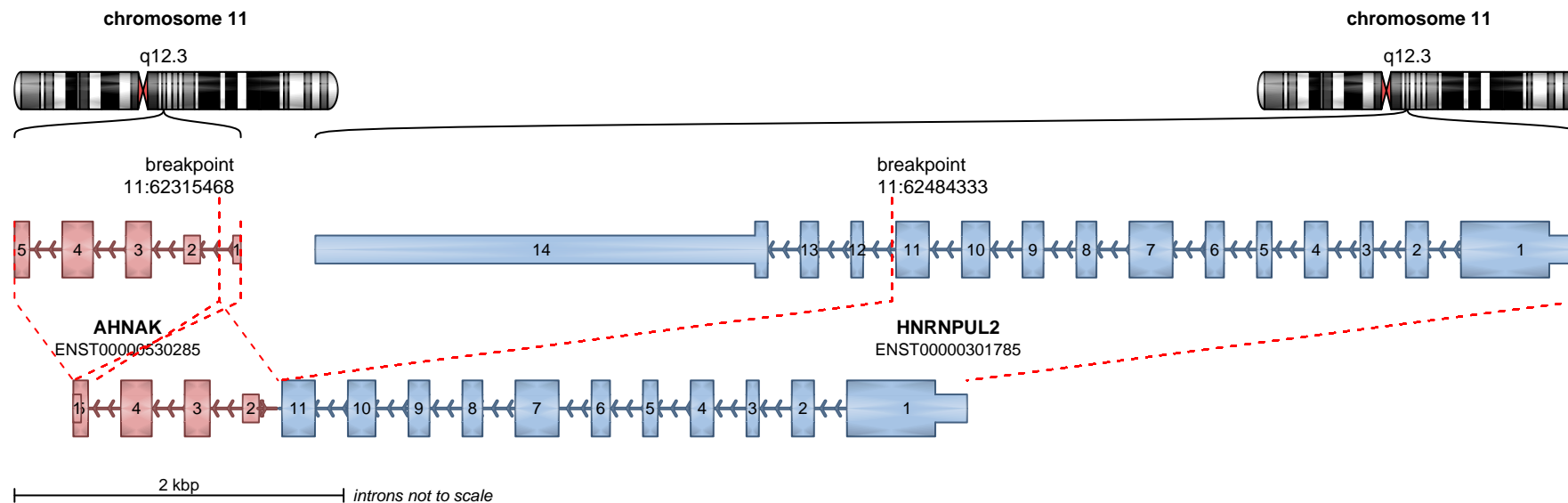
Total region count in KAZN = 15
 Total region count in HLA-DRB5 = 6
 Assembly score = _
 Fusion flag = 900



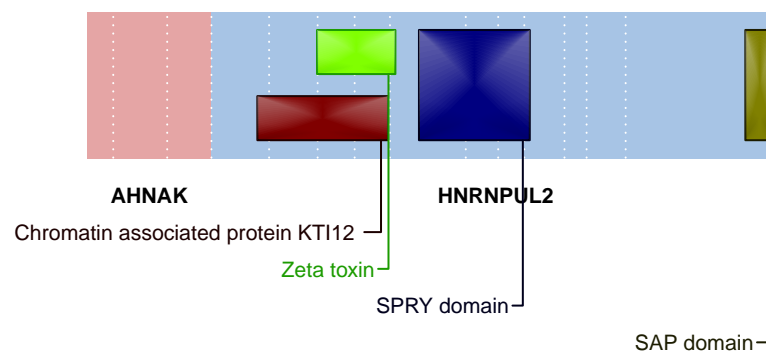


SUPPORTING READ COUNT

Total region count in PSEN2 = 13
 Total region count in SCAMP4 = 7
 Assembly score = 70
 Fusion flag = 820

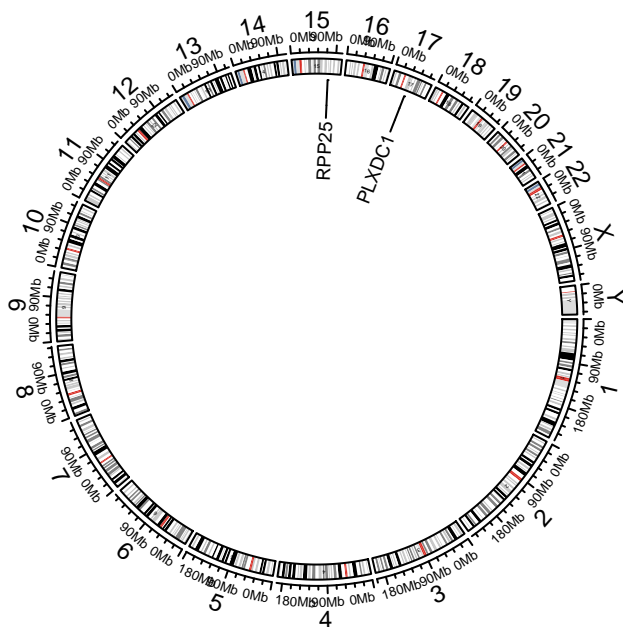
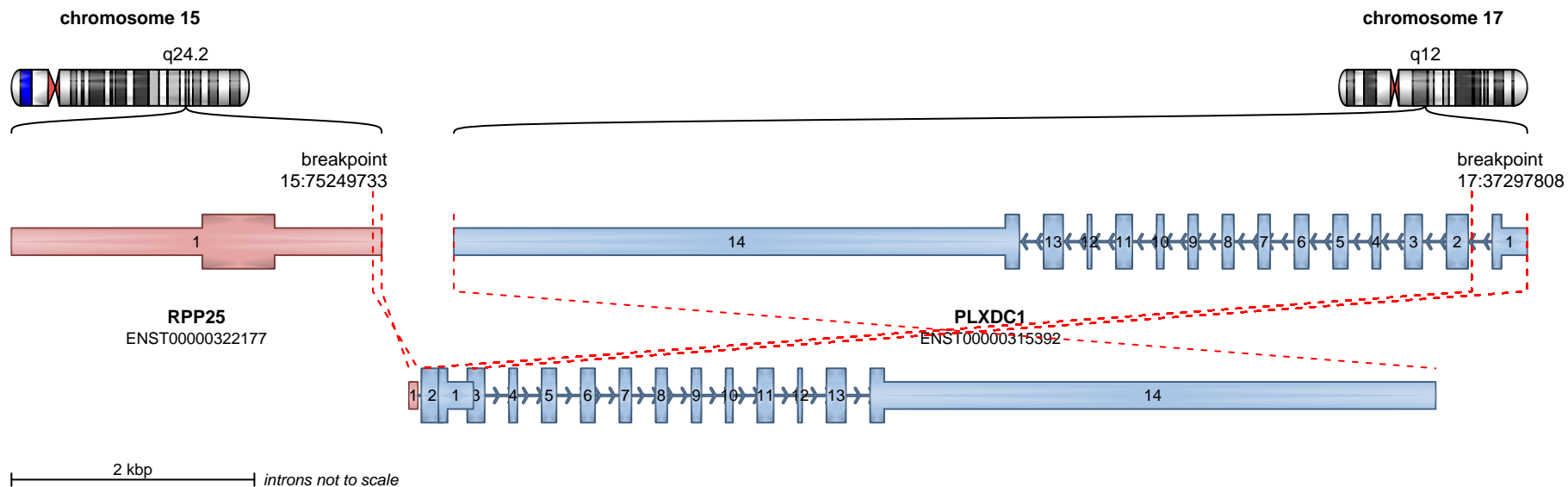


RETAINED PROTEIN DOMAINS



SUPPORTING READ COUNT

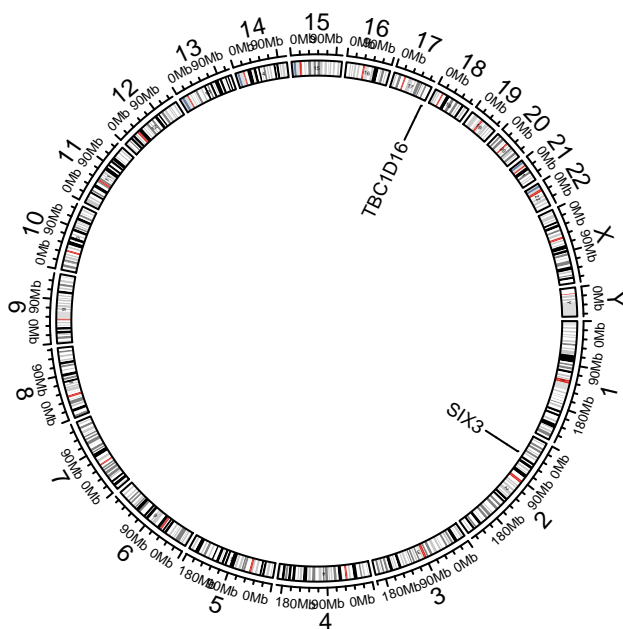
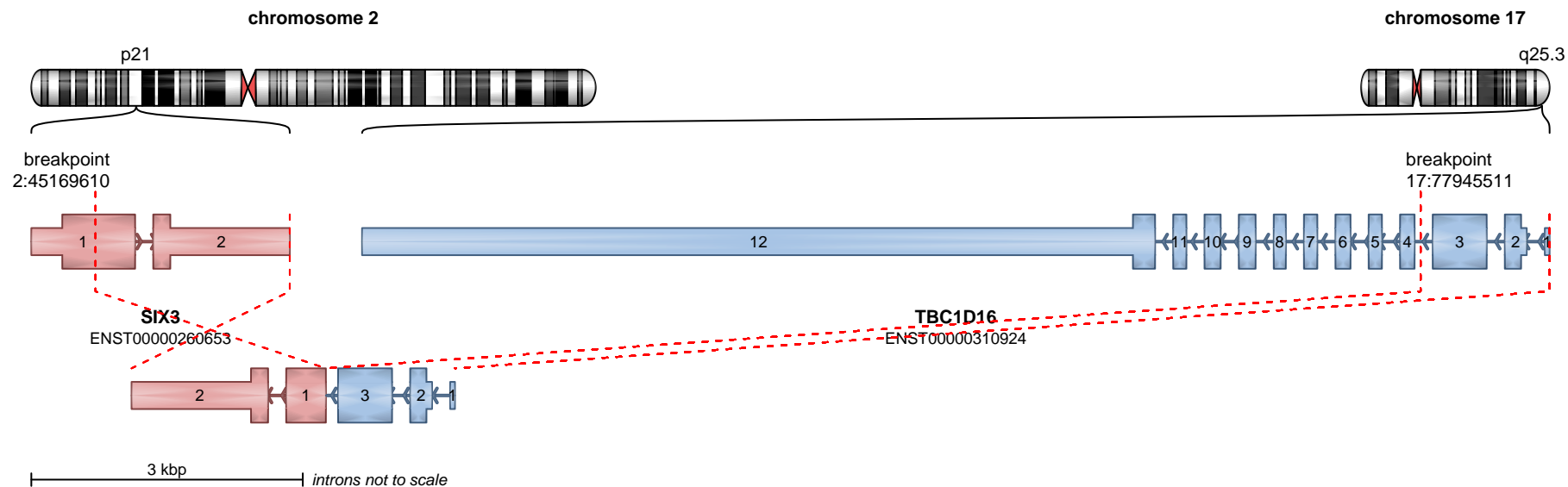
Total region count in AHNAK = 5
 Total region count in HNRNPUL2 = 14
 Assembly score = 100
 Fusion flag = 820



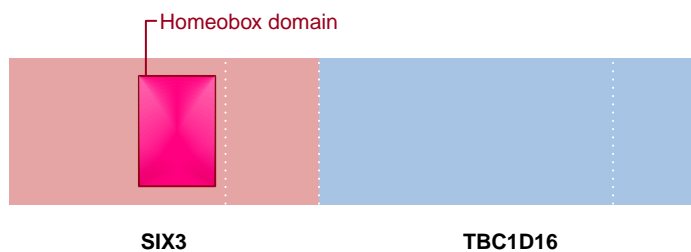
No protein domains retained in fusion.

SUPPORTING READ COUNT

Total region count in RPP25 = 1
Total region count in PLXDC1 = 14
Assembly score = 99
Fusion flag = 820

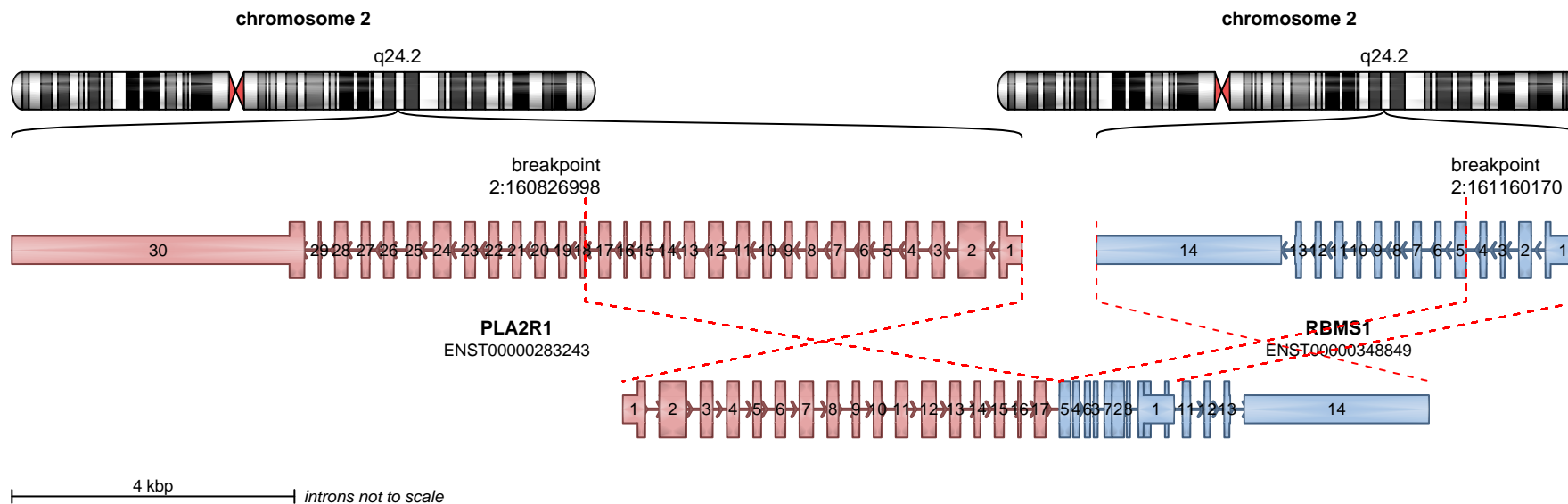


RETAINED PROTEIN DOMAINS

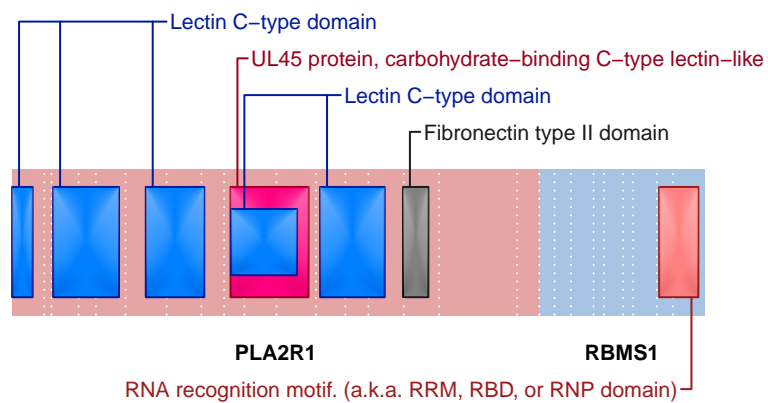


SUPPORTING READ COUNT

Total region count in SIX3 = 2
 Total region count in TBC1D16 = 12
 Assembly score = _
 Fusion flag = 740

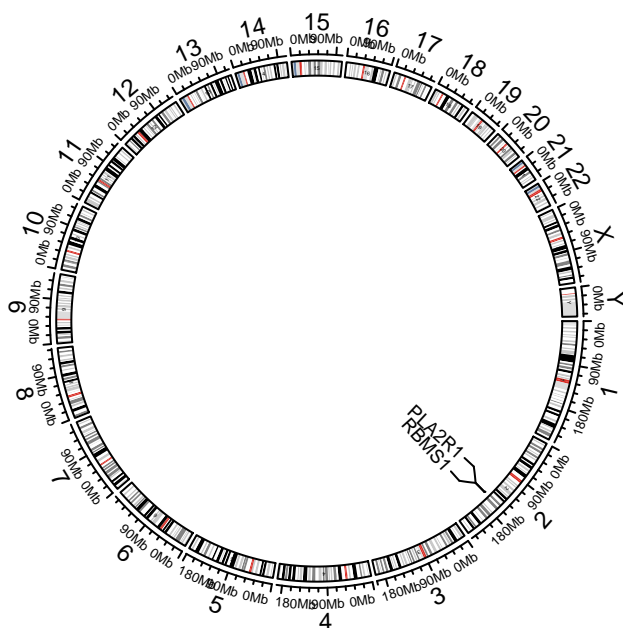


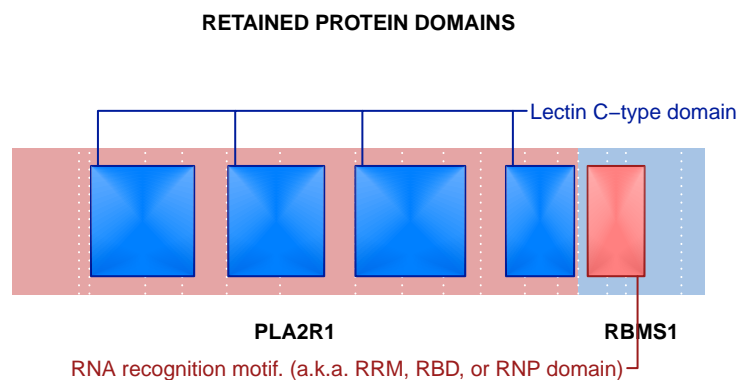
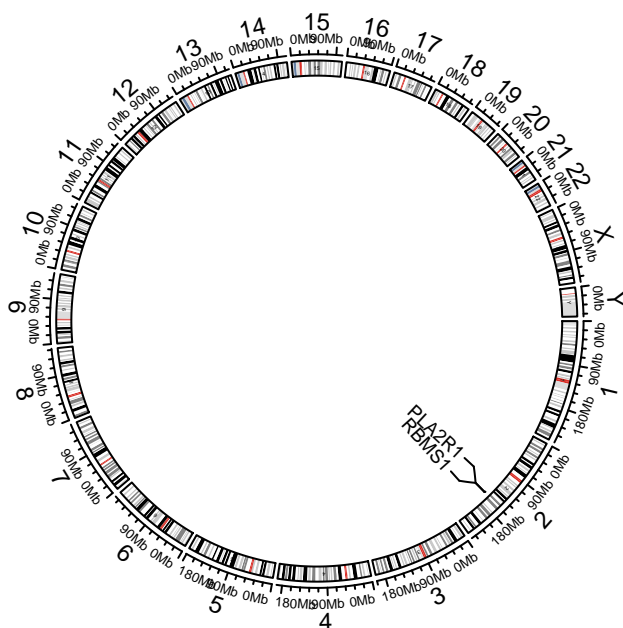
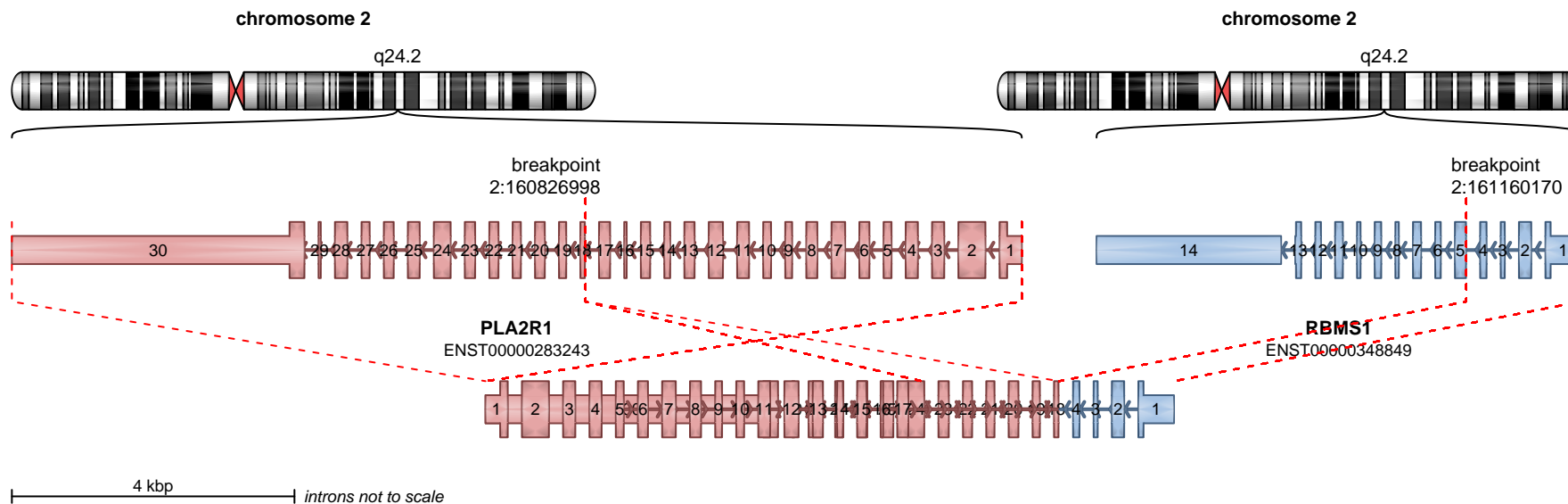
RETAINED PROTEIN DOMAINS



SUPPORTING READ COUNT

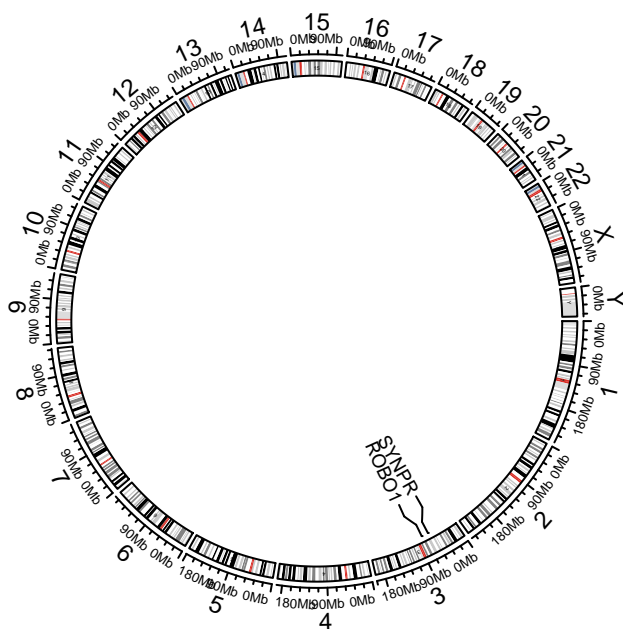
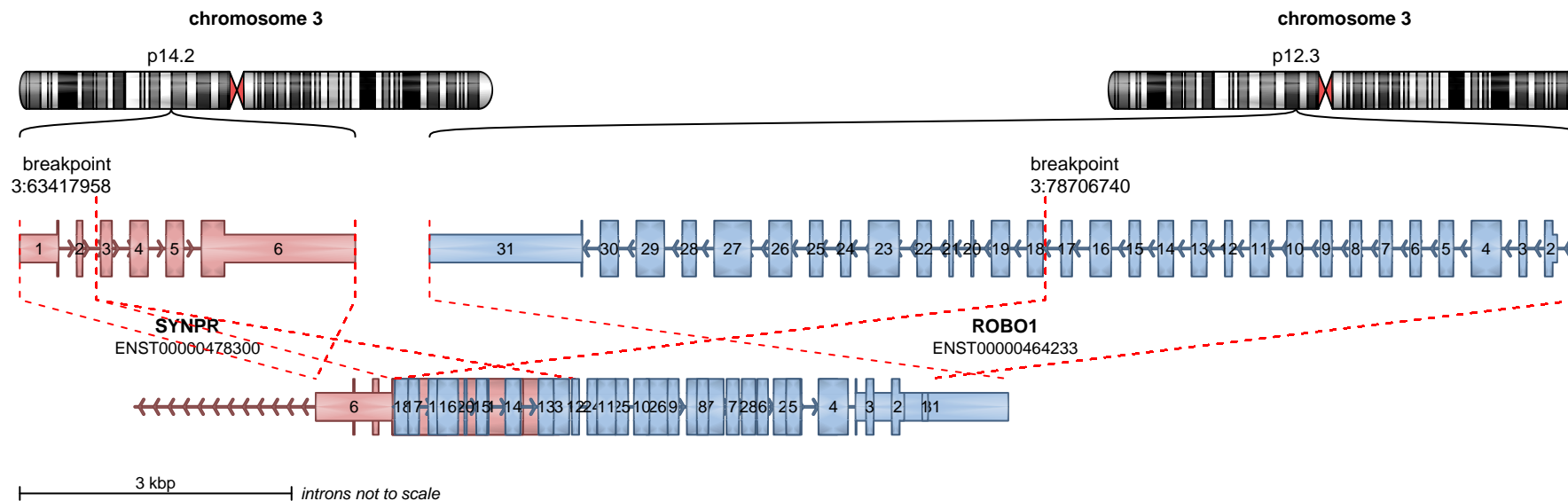
Total region count in PLA2R1 = 30
 Total region count in RBMS1 = 14
 Assembly score = 100
 Fusion flag = 900



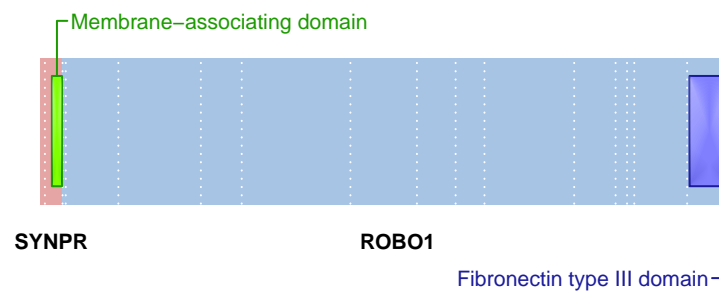


SUPPORTING READ COUNT

Total region count in PLA2R1 = 30
 Total region count in RBMS1 = 14
 Assembly score = 100
 Fusion flag = 900

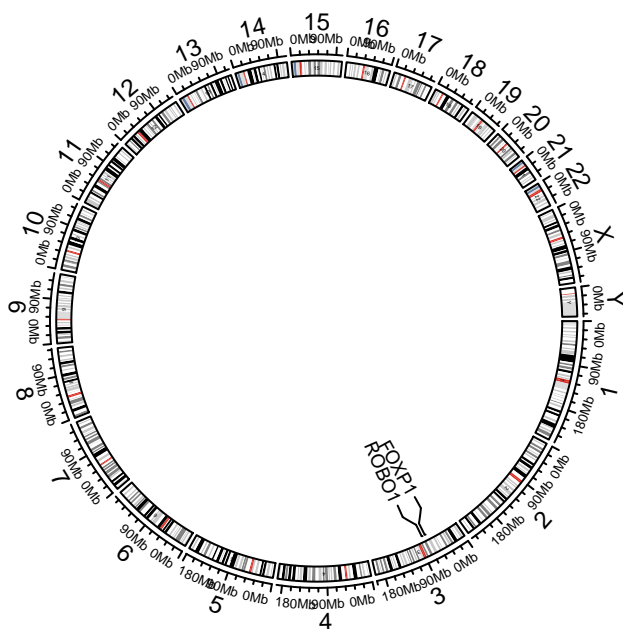
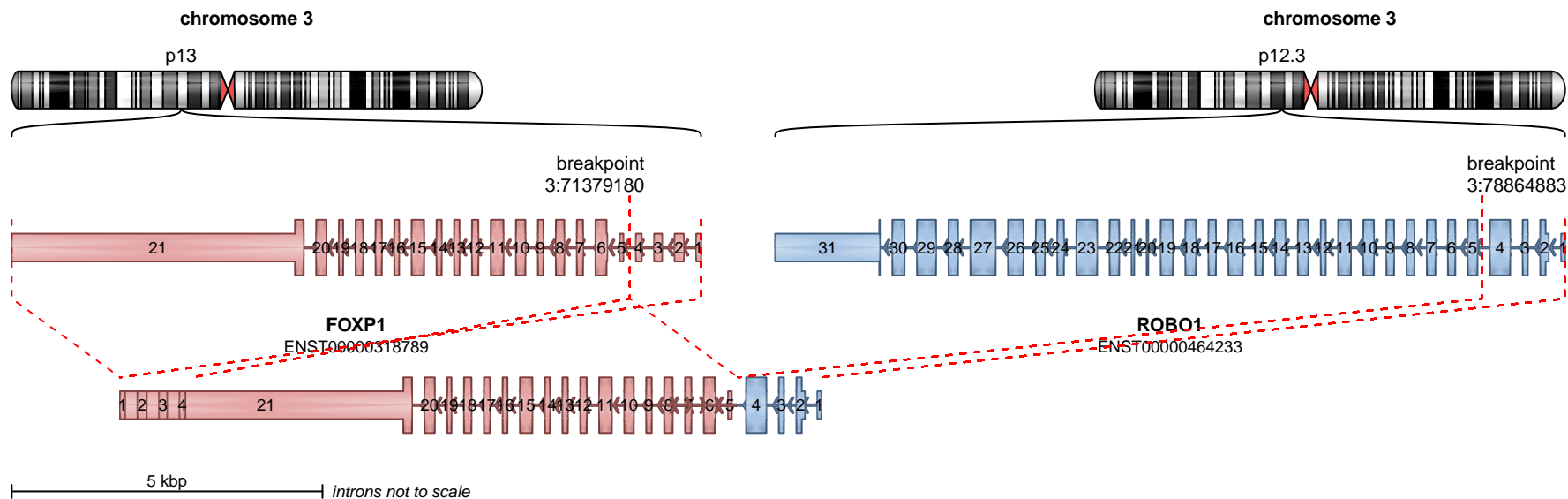


RETAINED PROTEIN DOMAINS

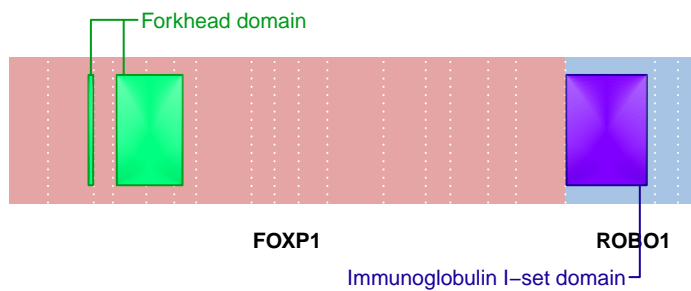


SUPPORTING READ COUNT

Total region count in SYNPR = 6
 Total region count in ROBO1 = 31
 Assembly score = 100
 Fusion flag = 900

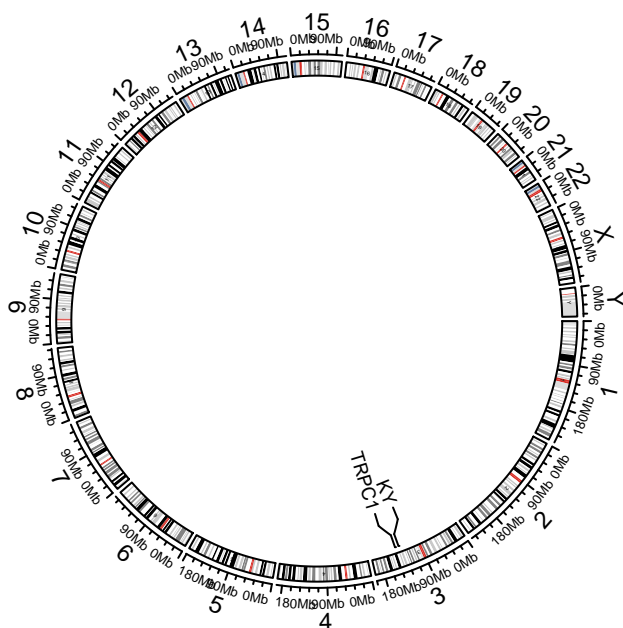
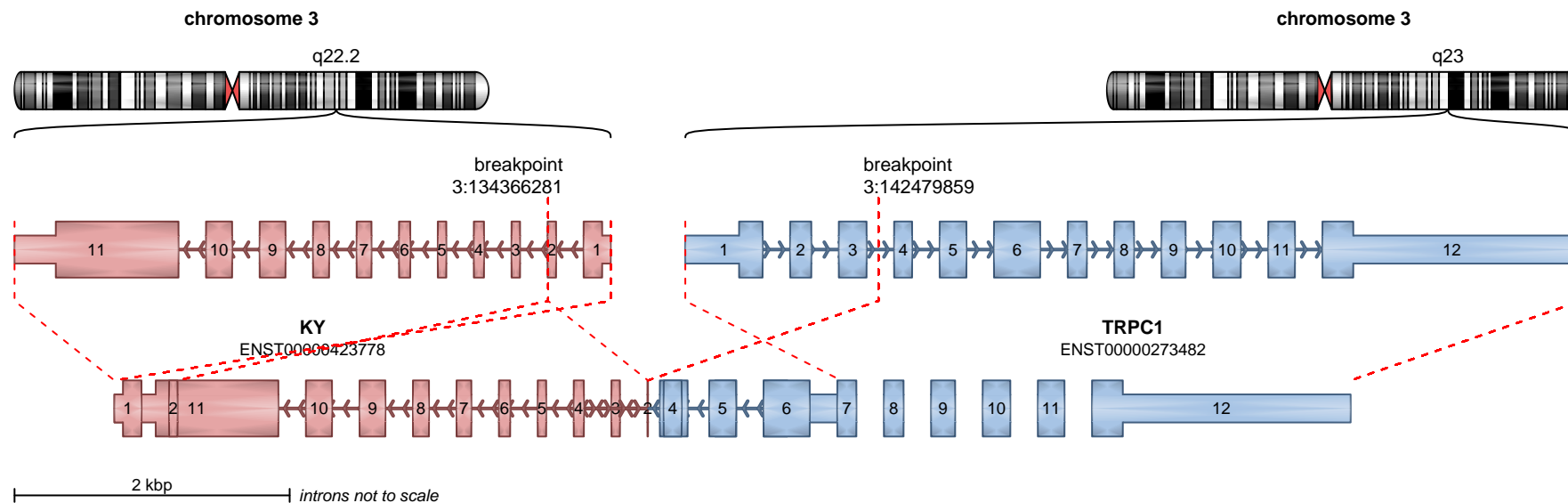


RETAINED PROTEIN DOMAINS

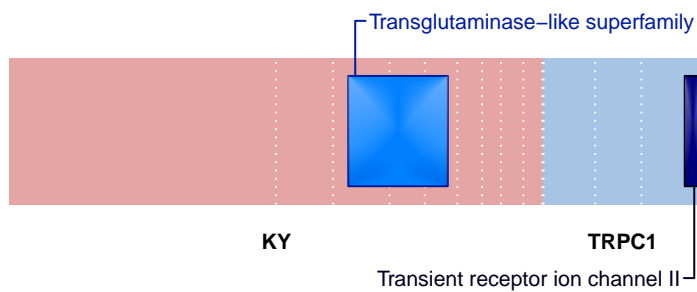


SUPPORTING READ COUNT

Total region count in FOXP1 = 21
 Total region count in ROBO1 = 31
 Assembly score = 99
 Fusion flag = 820

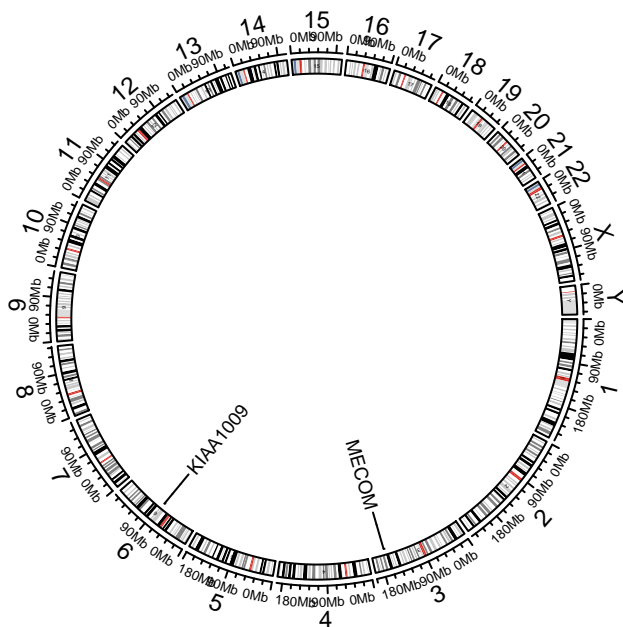
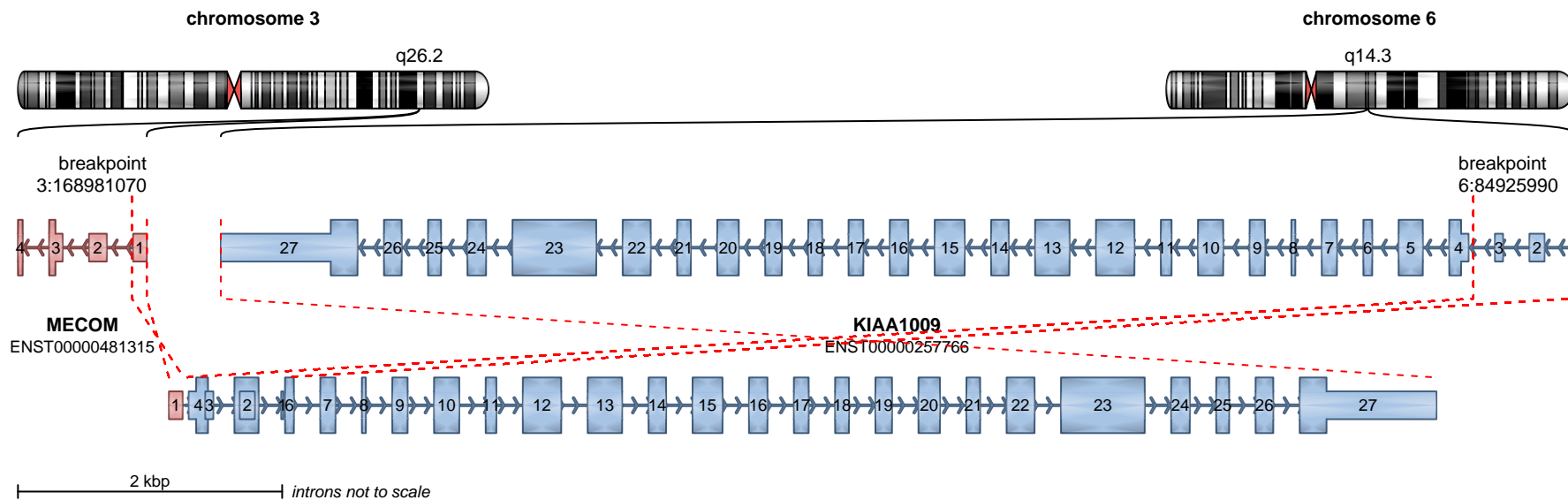


RETAINED PROTEIN DOMAINS



SUPPORTING READ COUNT

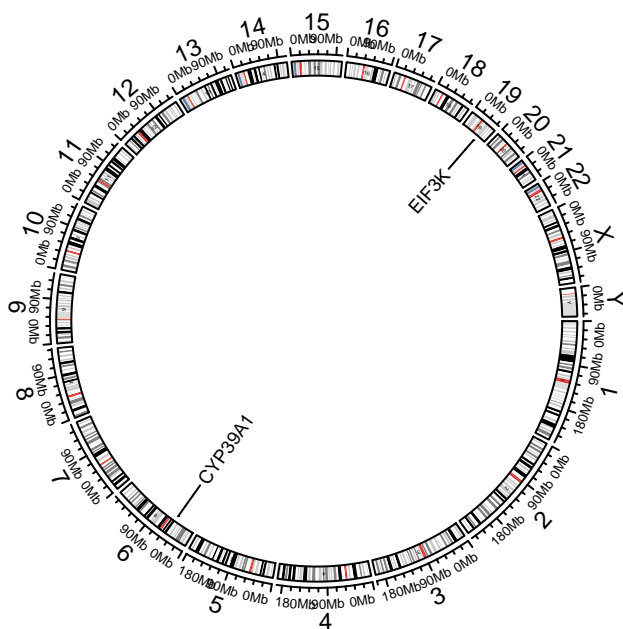
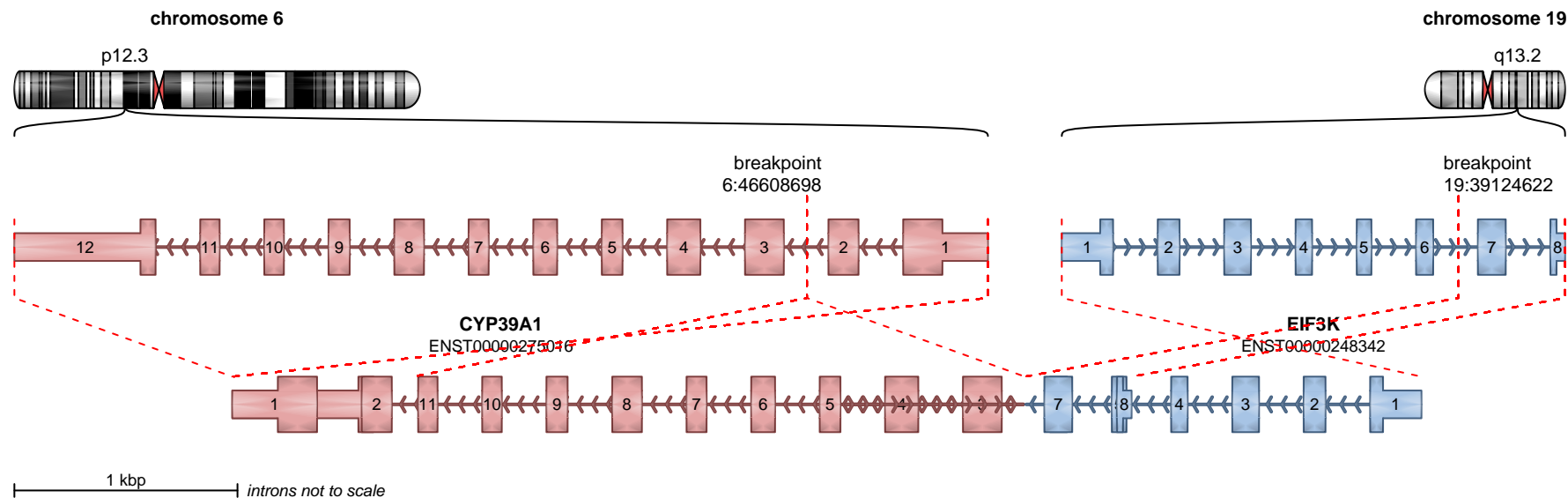
Total region count in KY = 11
 Total region count in TRPC1 = 12
 Assembly score = 100
 Fusion flag = 740



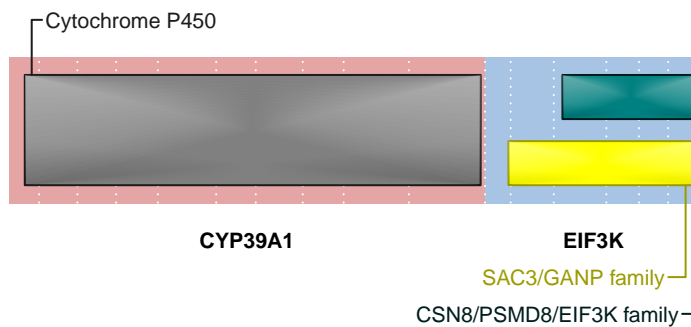
No protein domains retained in fusion.

SUPPORTING READ COUNT

Total region count in MECOM = 4
 Total region count in KIAA1009 = 27
 Assembly score = 100
 Fusion flag = 860

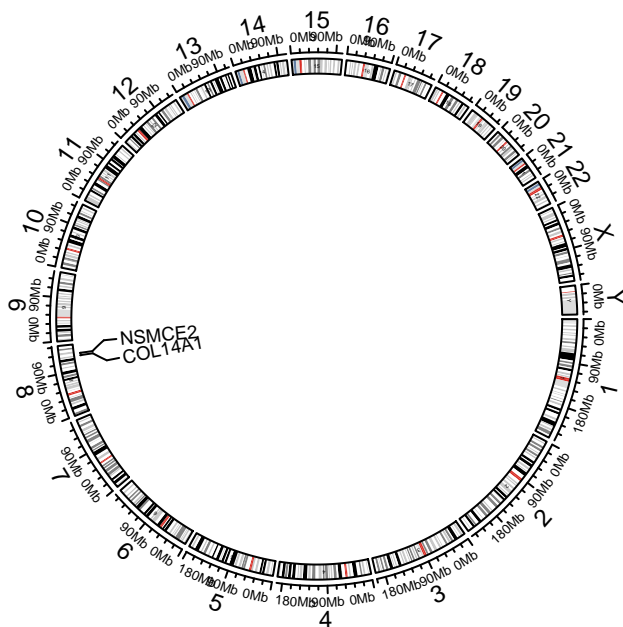
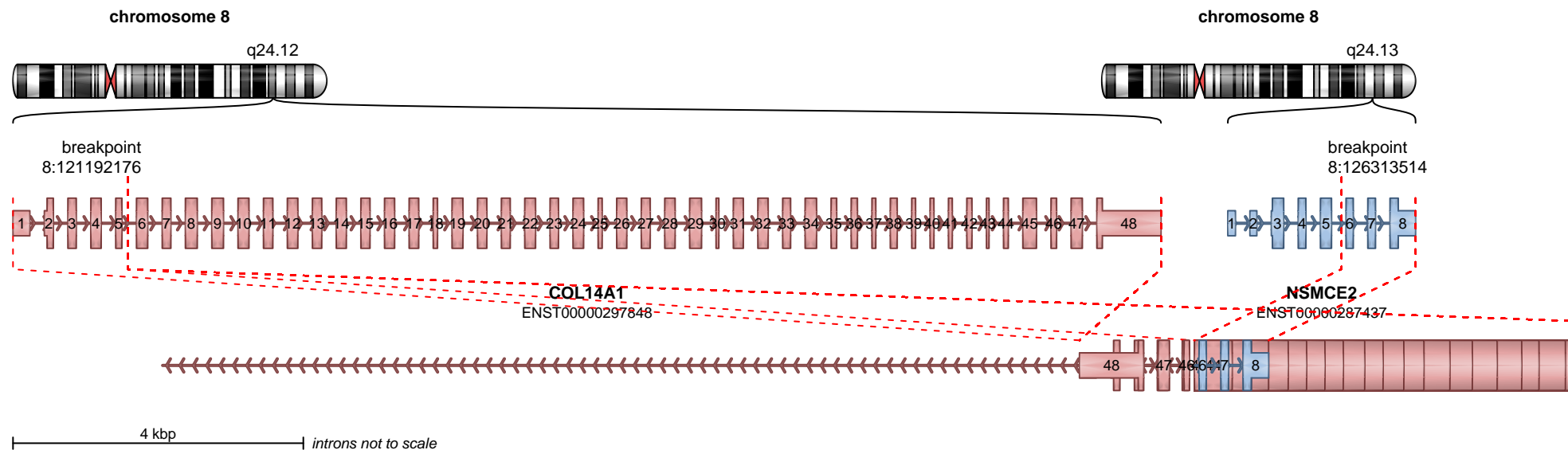


RETAINED PROTEIN DOMAINS

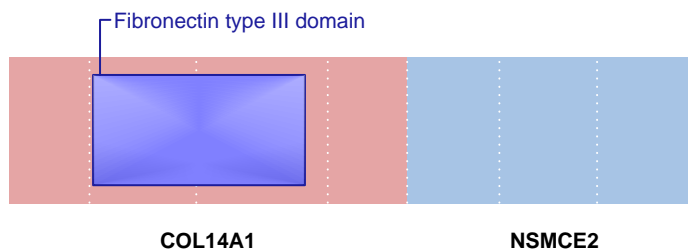


SUPPORTING READ COUNT

Total region count in CYP39A1 = 12
 Total region count in EIF3K = 8
 Assembly score = 99
 Fusion flag = 900



RETAINED PROTEIN DOMAINS



SUPPORTING READ COUNT

Total region count in COL14A1 = 48
 Total region count in NSMCE2 = 8
 Assembly score = 99
 Fusion flag = 900