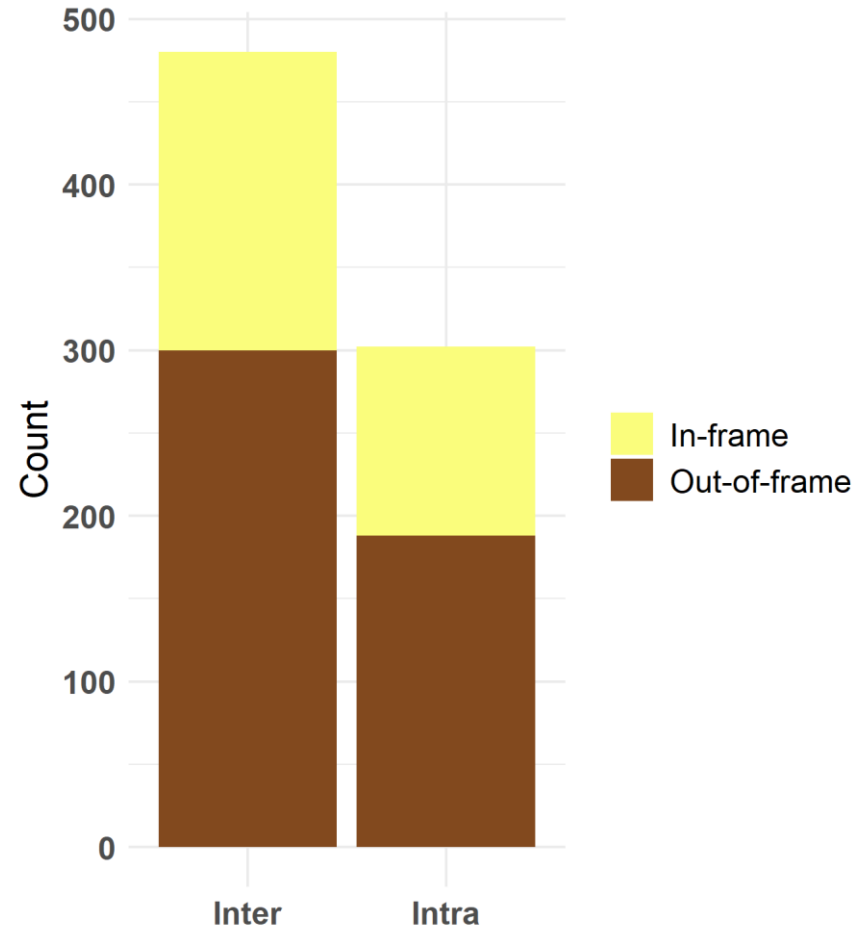
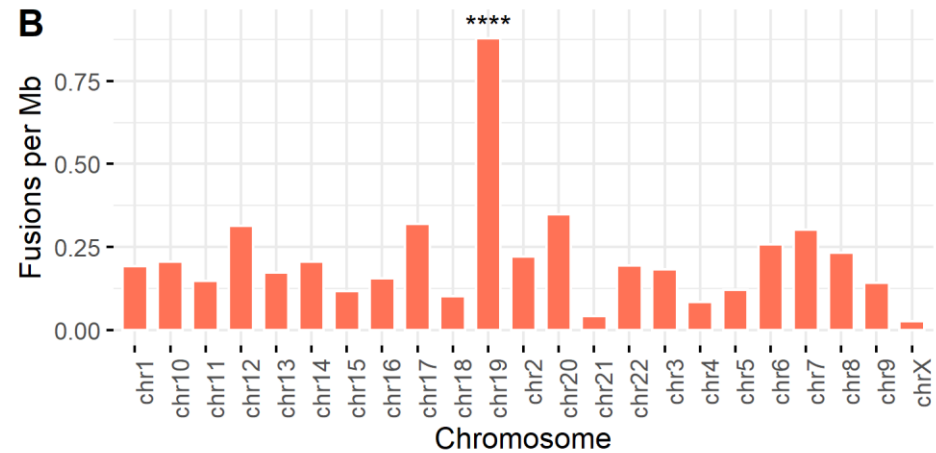
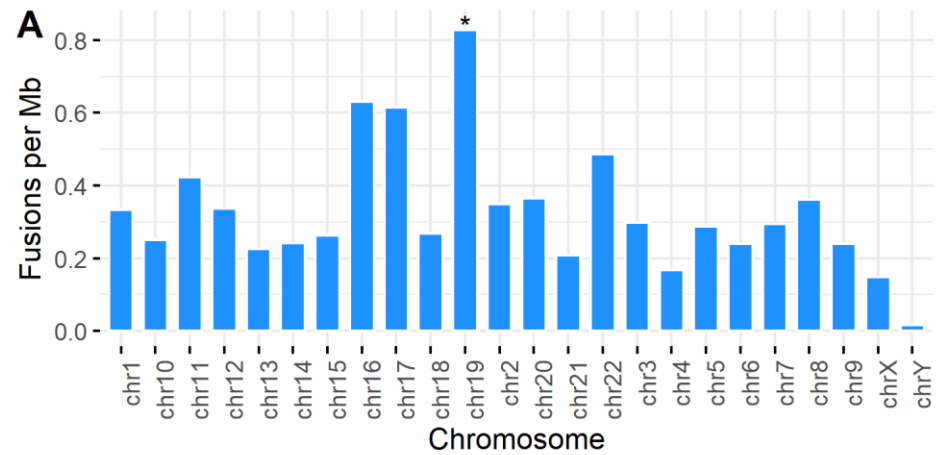


Supplementary figure 1. Classification of expressed features (genes) detected across all samples using RNA-sequencing. The biotypes were obtained with biomaRt Bioconductor package.



Supplementary figure 2. Inter- and intra-chromosomal transcript fusions categorized by their ability produce viable protein based on in- or out-of-frame status.



Supplementary figure 3.  
Distributions of inter – (A) and  
intra-chromosomal (B) fusions  
across chromosomes, normalized  
per megabase. Chromosome 19 is  
a significant outlier according  
Grubb's test (A:  $p < 0.05$ ; B:  
 $p < 0.0001$ )