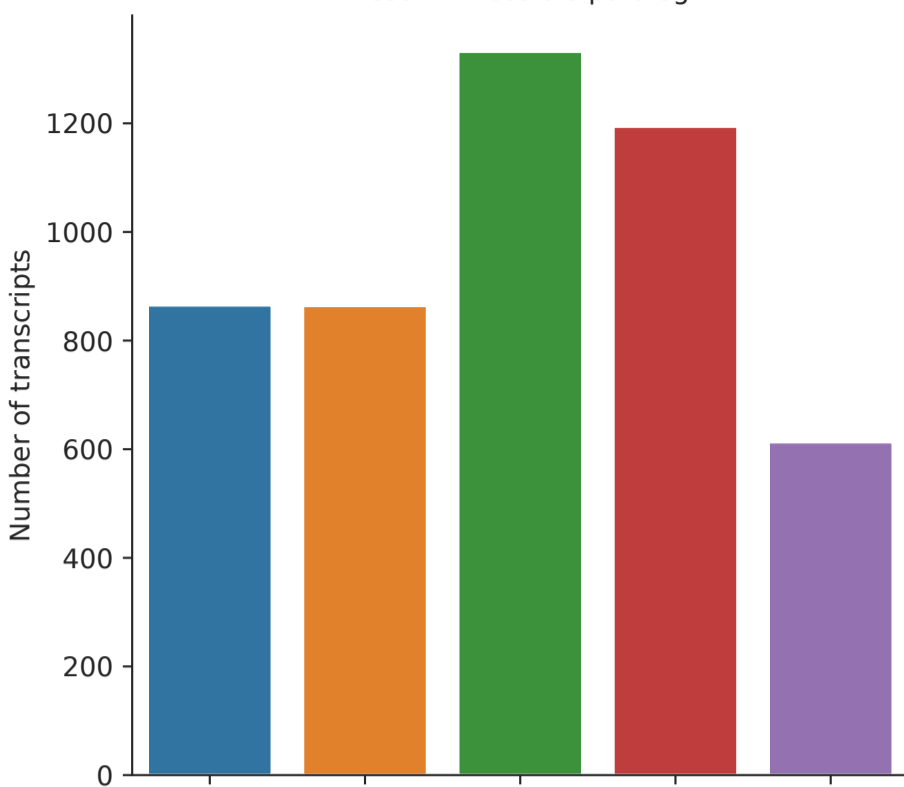
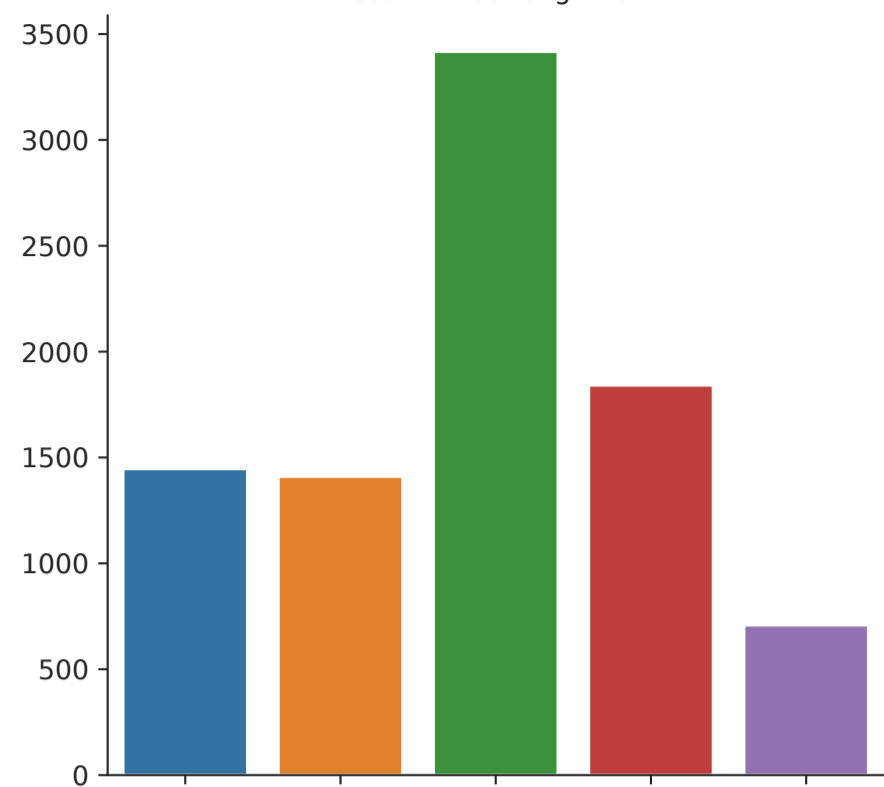


# Incorporation of de-novo predictions

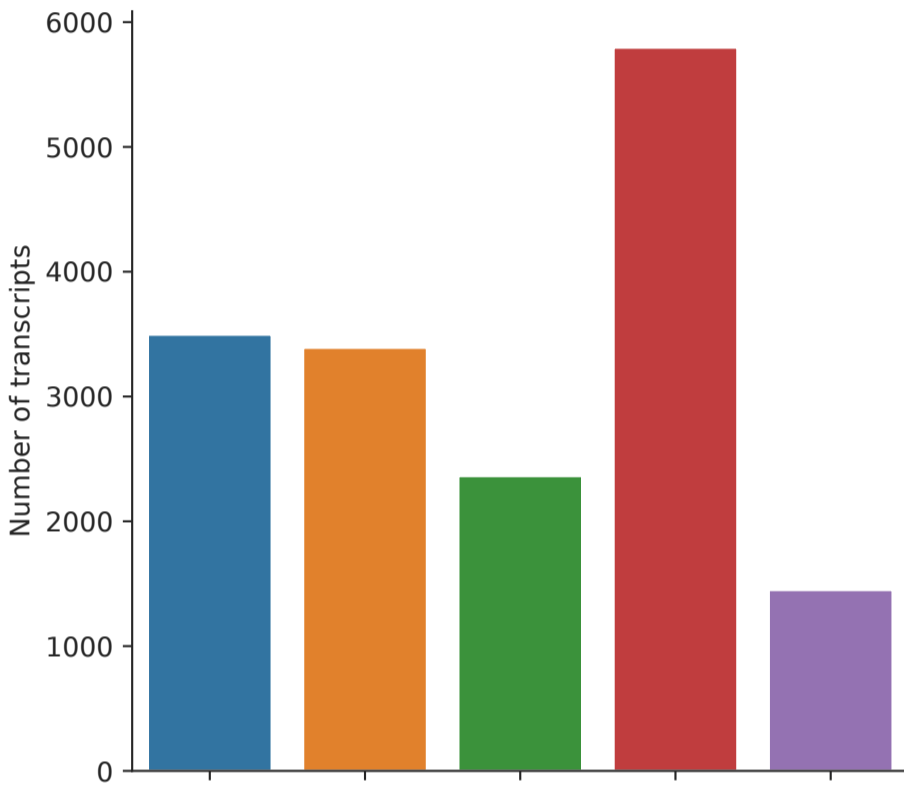
### Result = Possible paralog



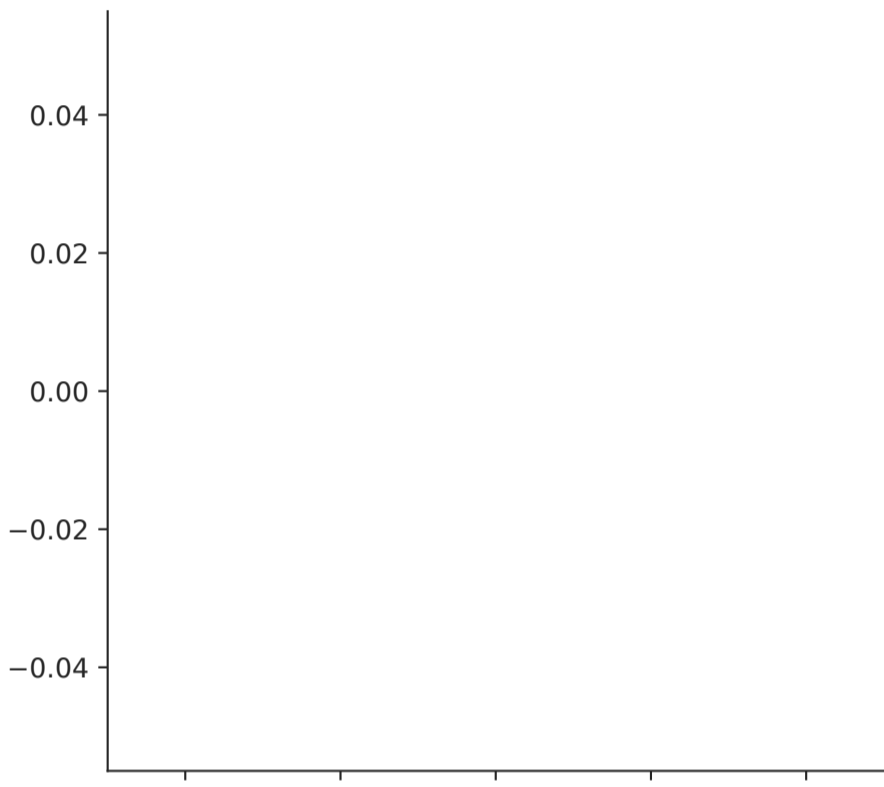
### Result = Poor alignment



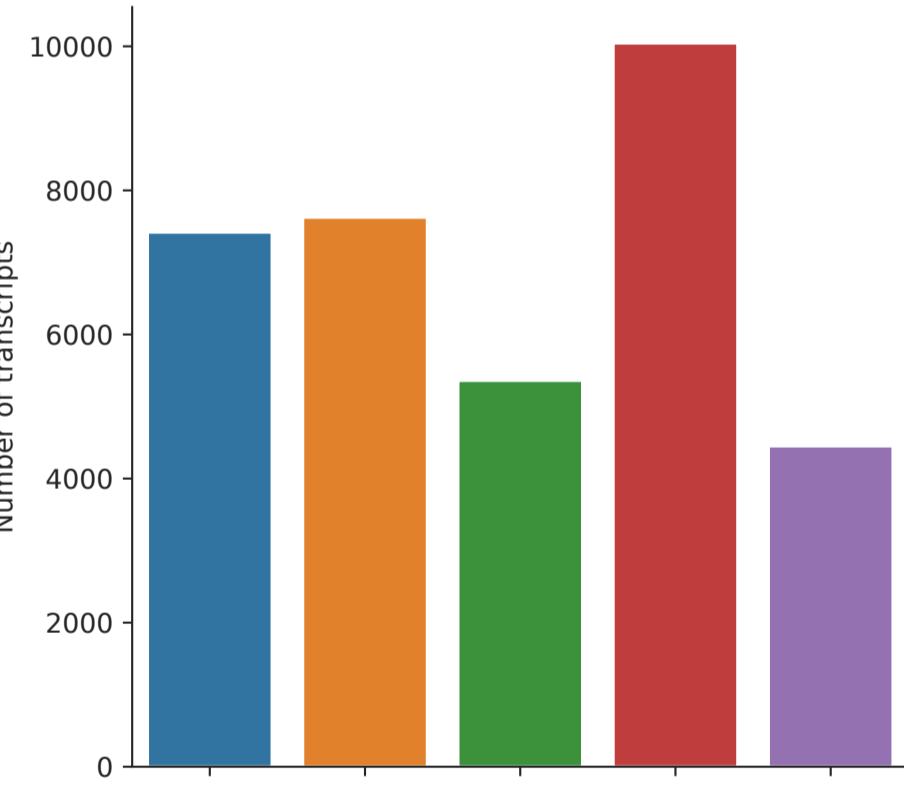
### Result = Putative novel



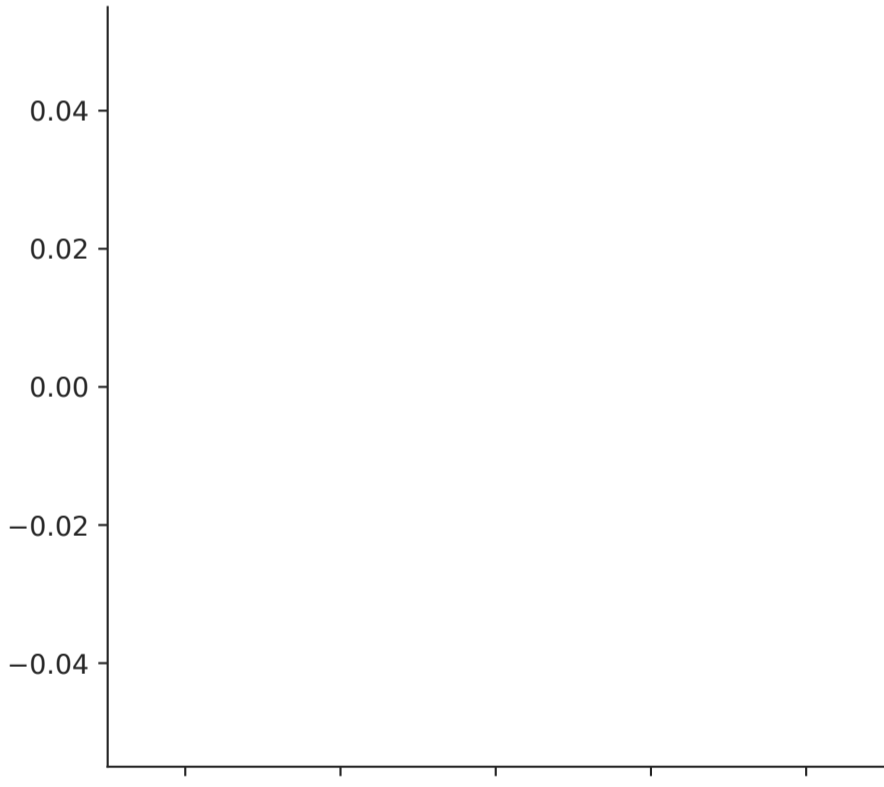
### Result = Possible fusion



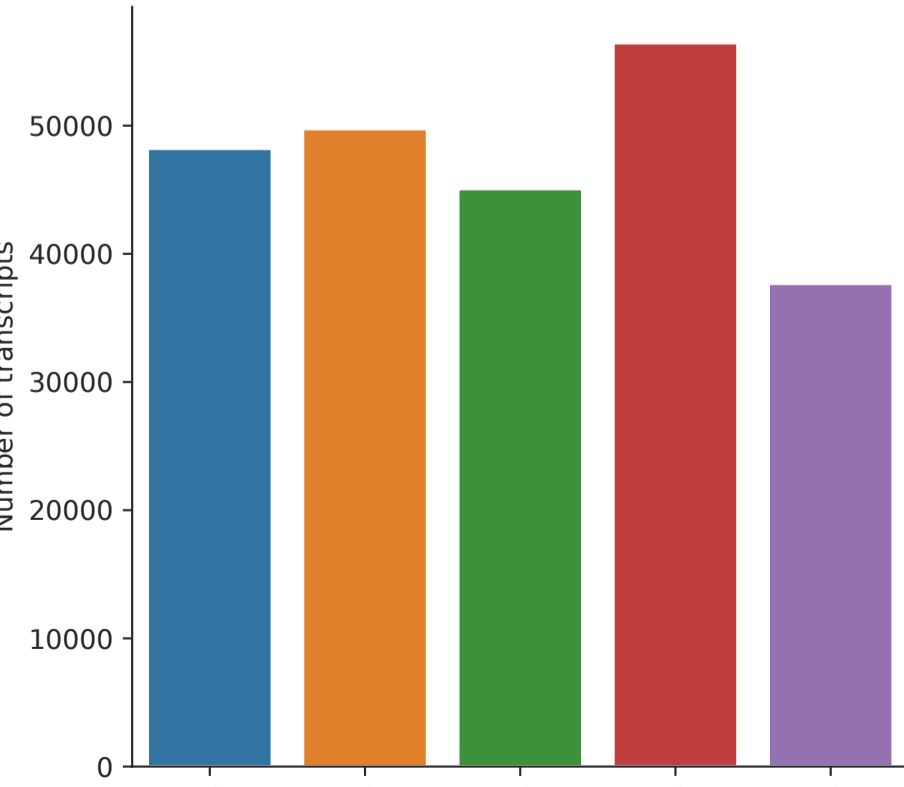
### Result = Putative novel isoform



### Result = Single exon



### Result = Discarded



GCF\_029289425.1  
GCF\_028858775.1  
GCF\_029281585.1  
GCF\_028885625.1  
GCF\_028885655.1  
genome

GCF\_029289425.1  
GCF\_028858775.1  
GCF\_029281585.1  
GCF\_028885625.1  
GCF\_028885655.1  
genome