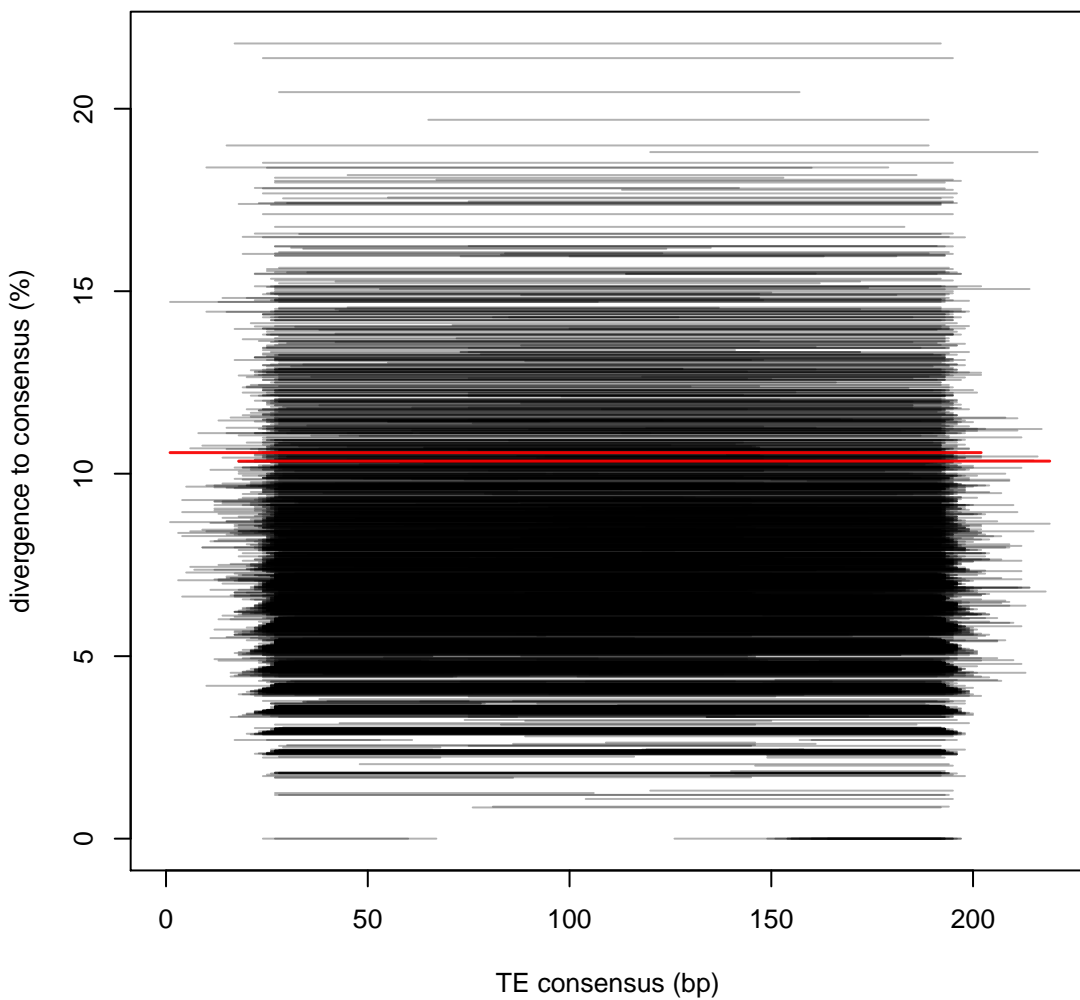
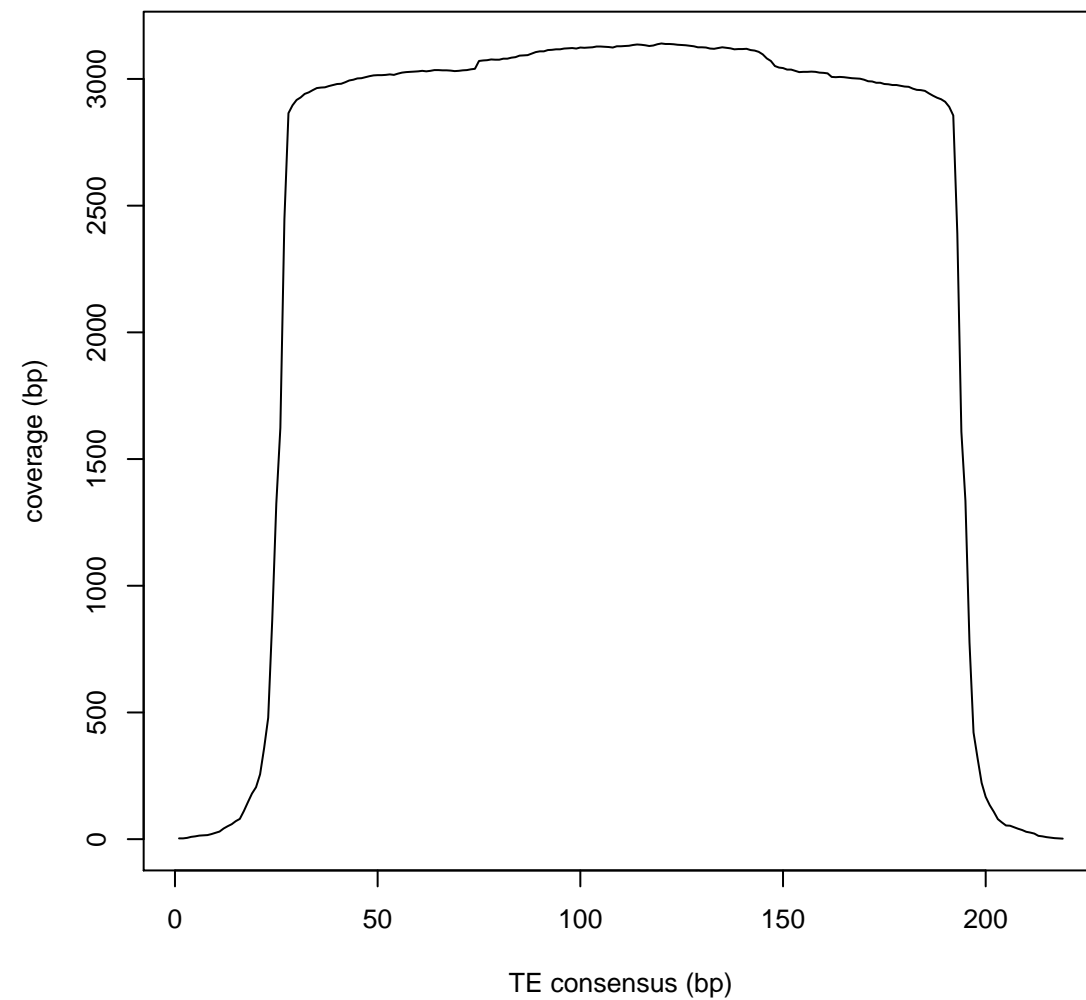


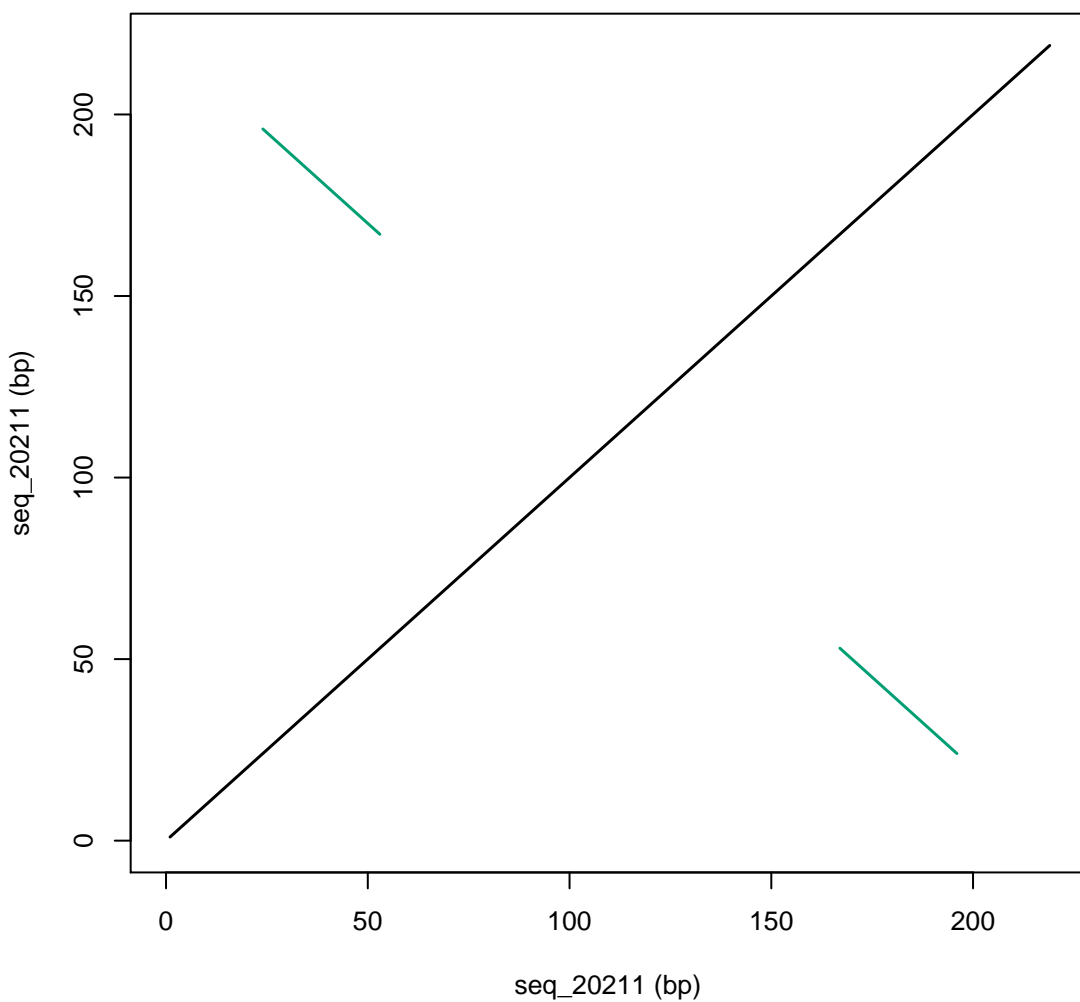
TE: seq\_20211  
consensus size: 219bp; fragments: 3260; full length: 2 ( $\geq 197.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

