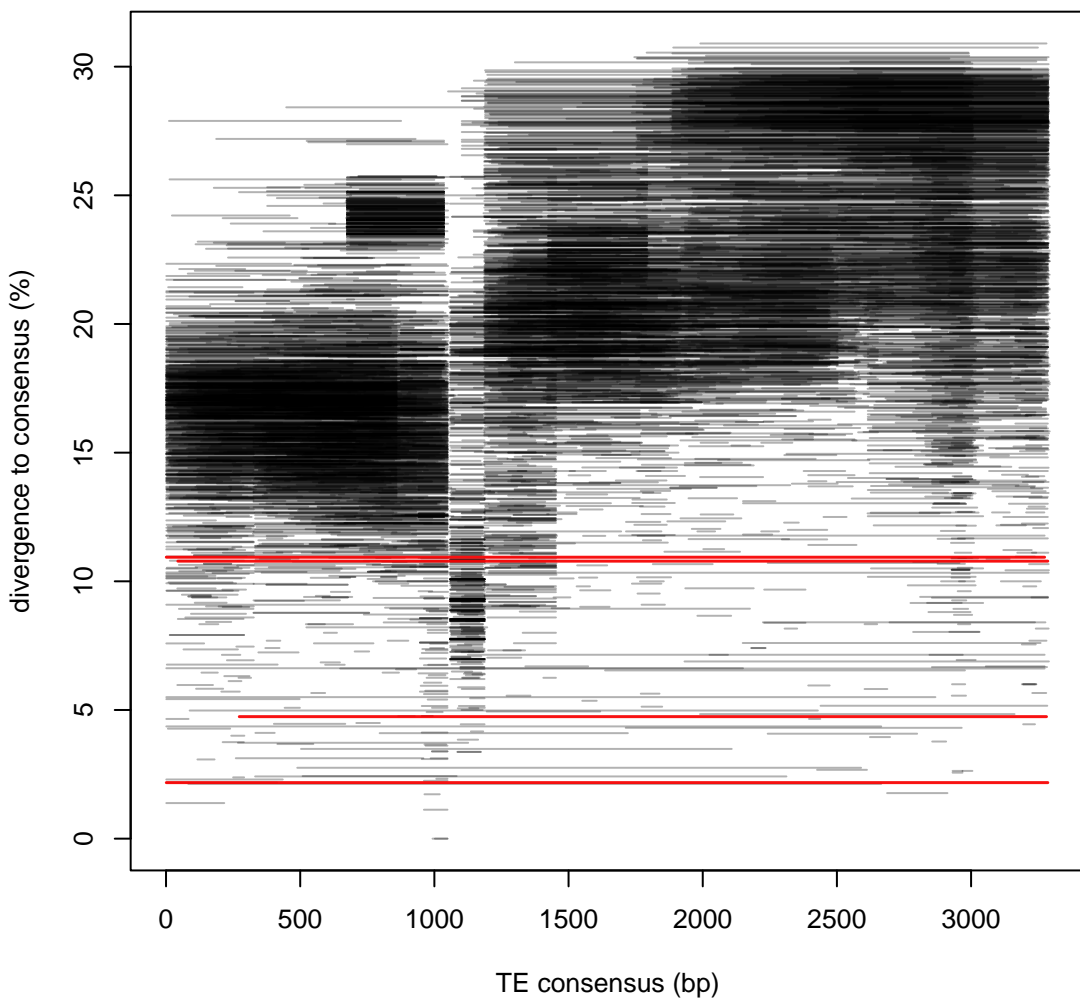
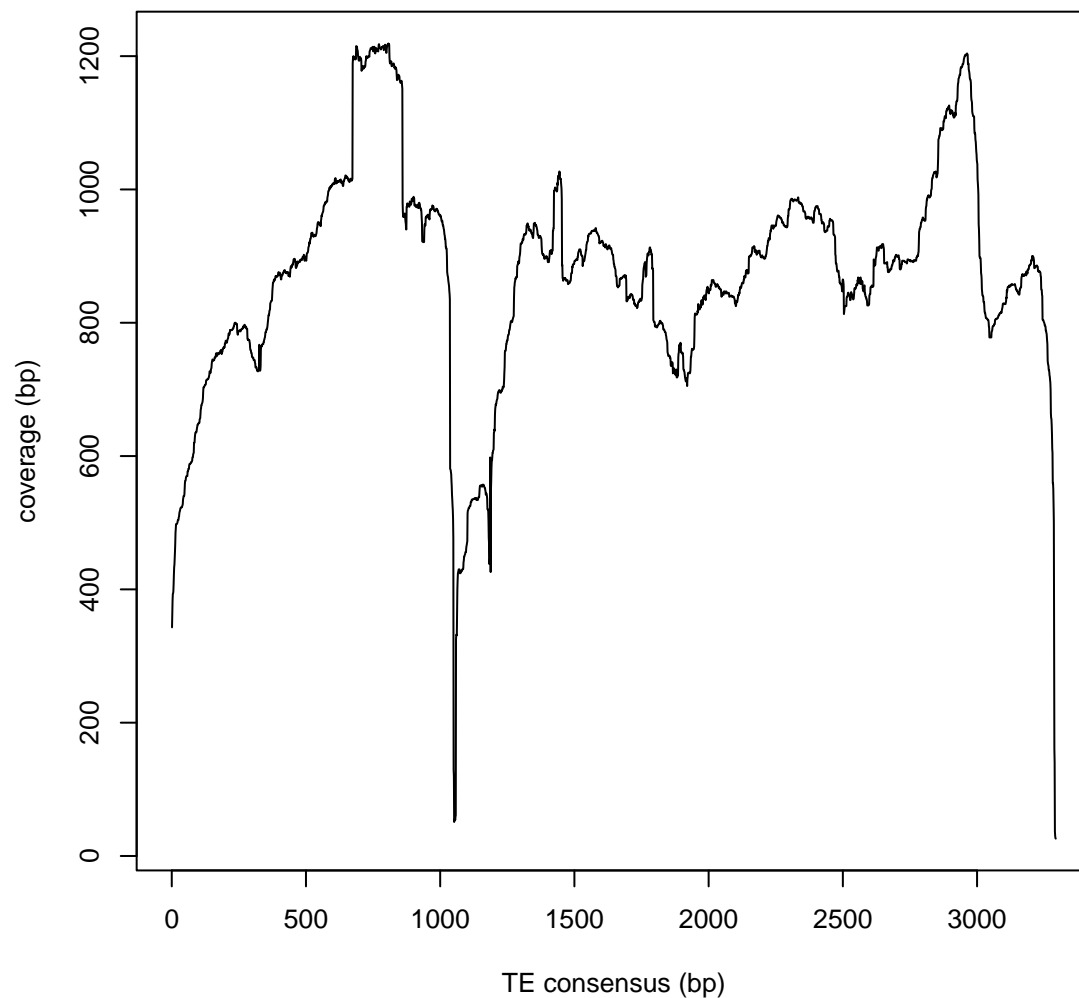


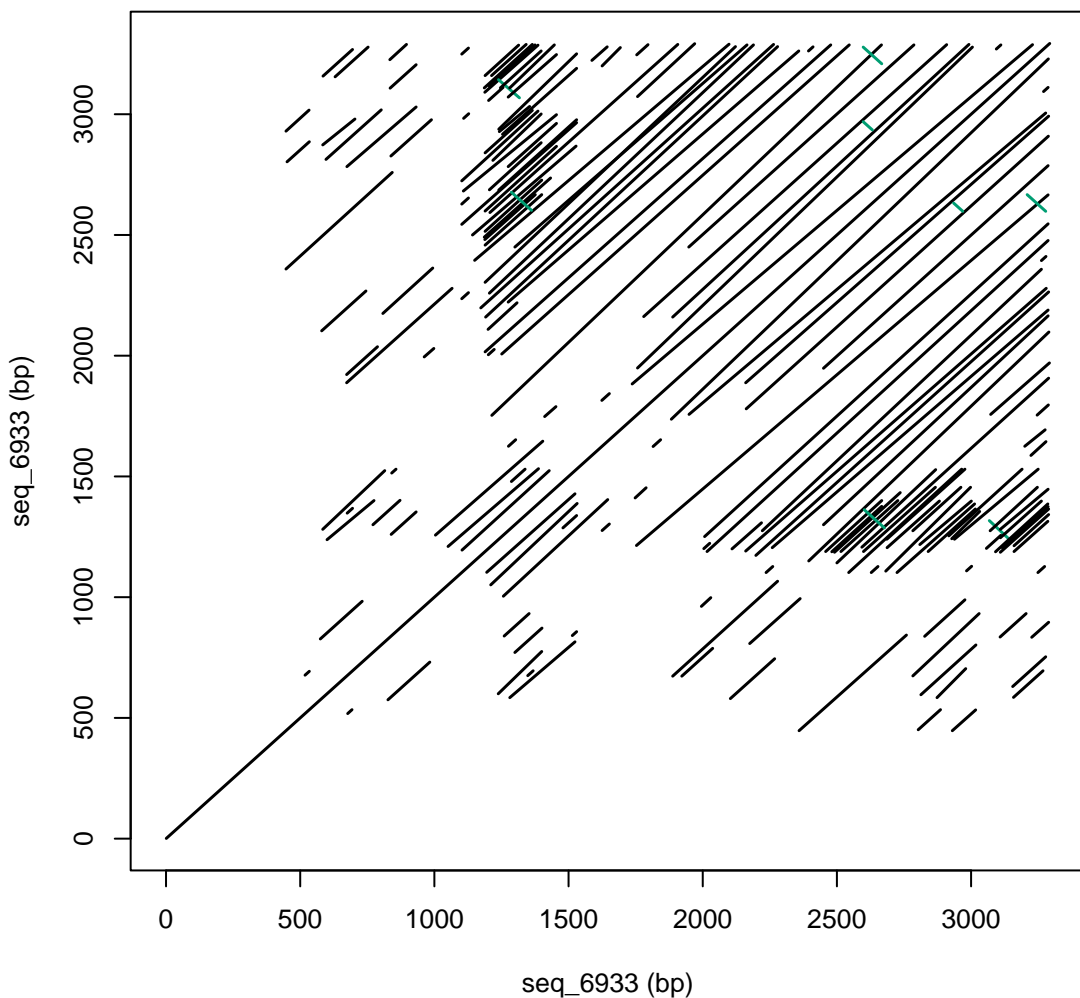
TE: seq\_6933  
consensus size: 3293bp; fragments: 5648; full length: 4 ( $\geq 2963.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

