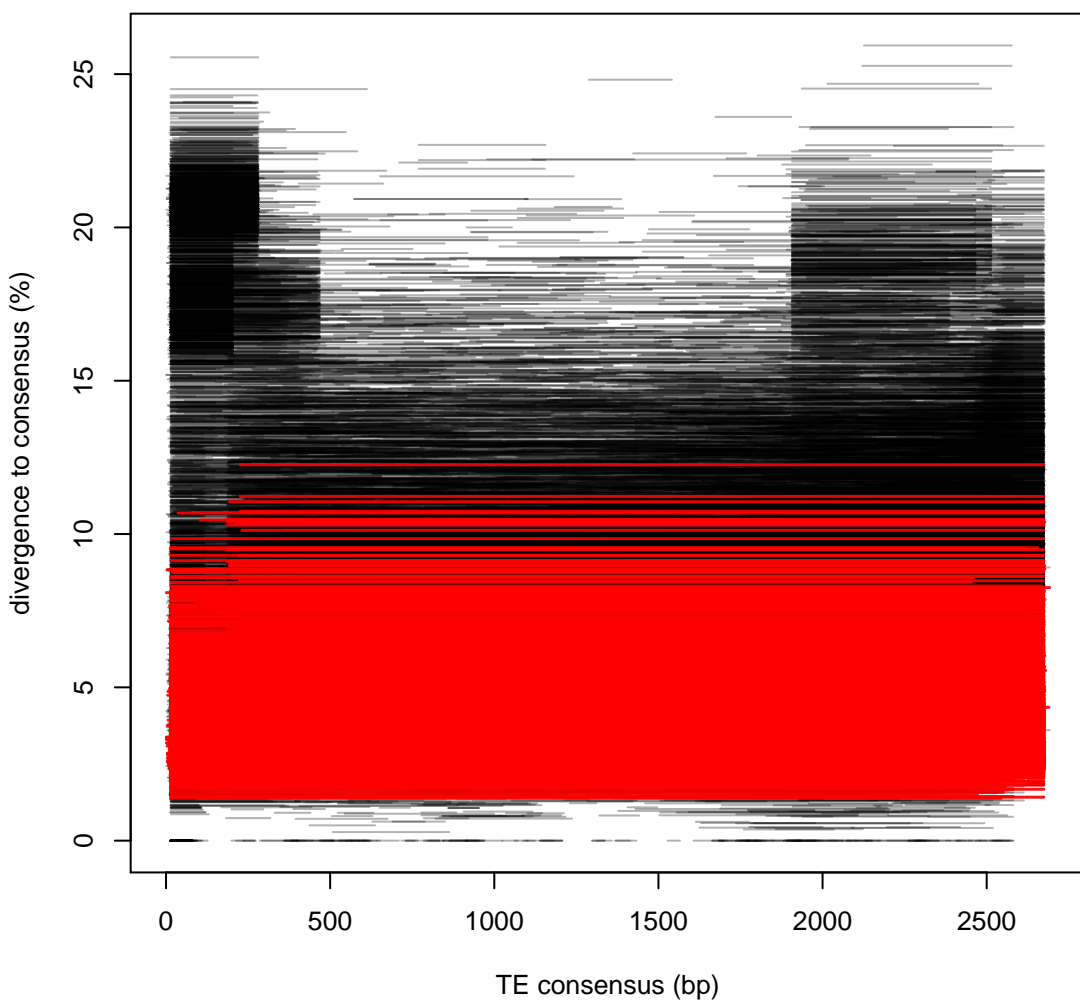
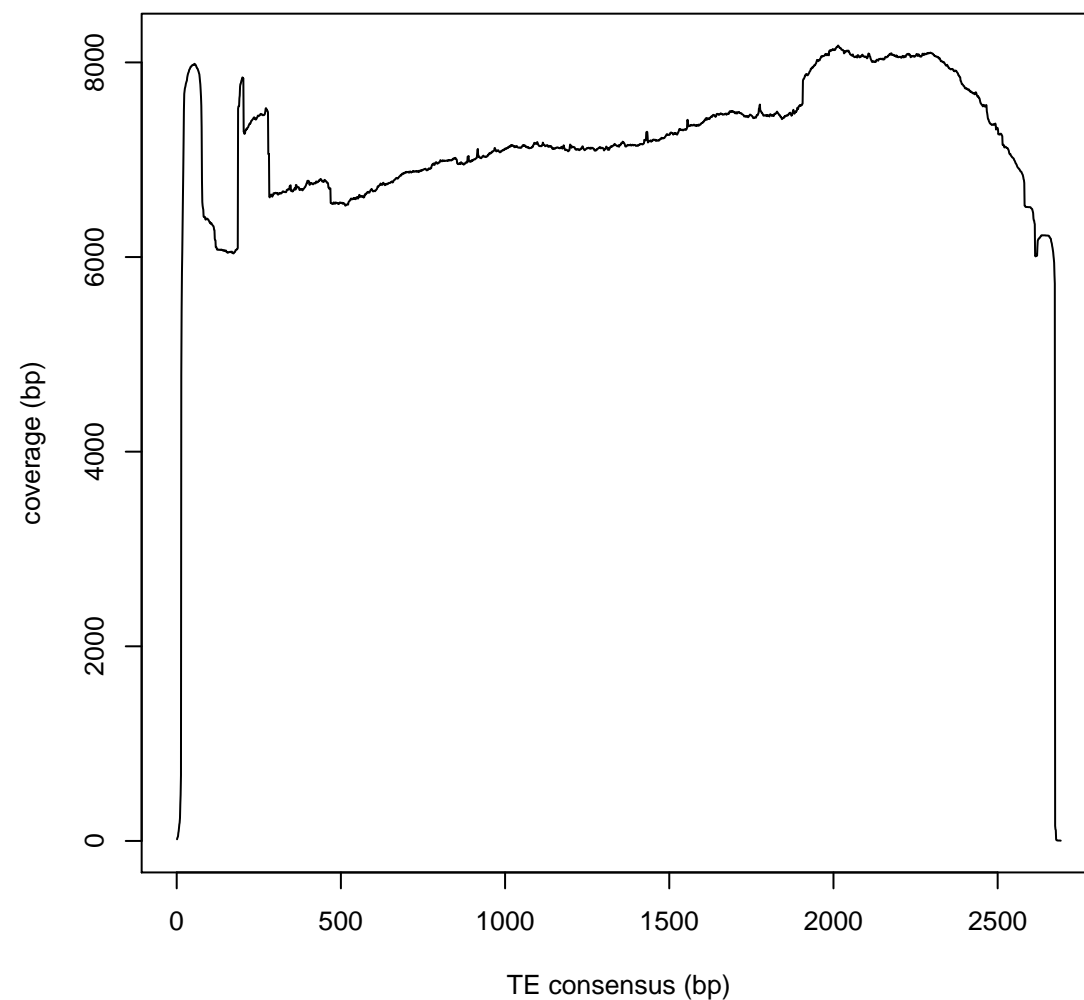


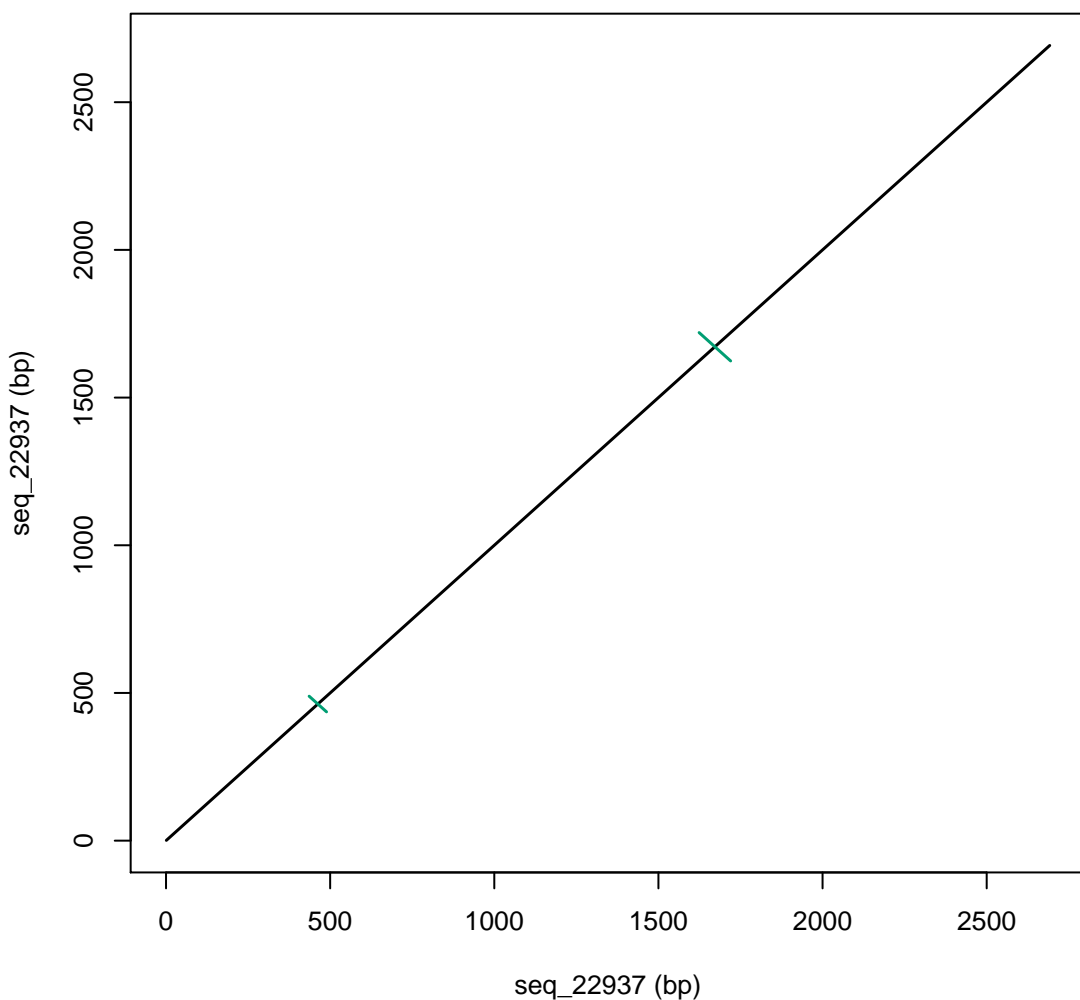
TE: seq\_22937  
consensus size: 2692bp; fragments: 26208; full length: 2167 ( $\geq 2422.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

