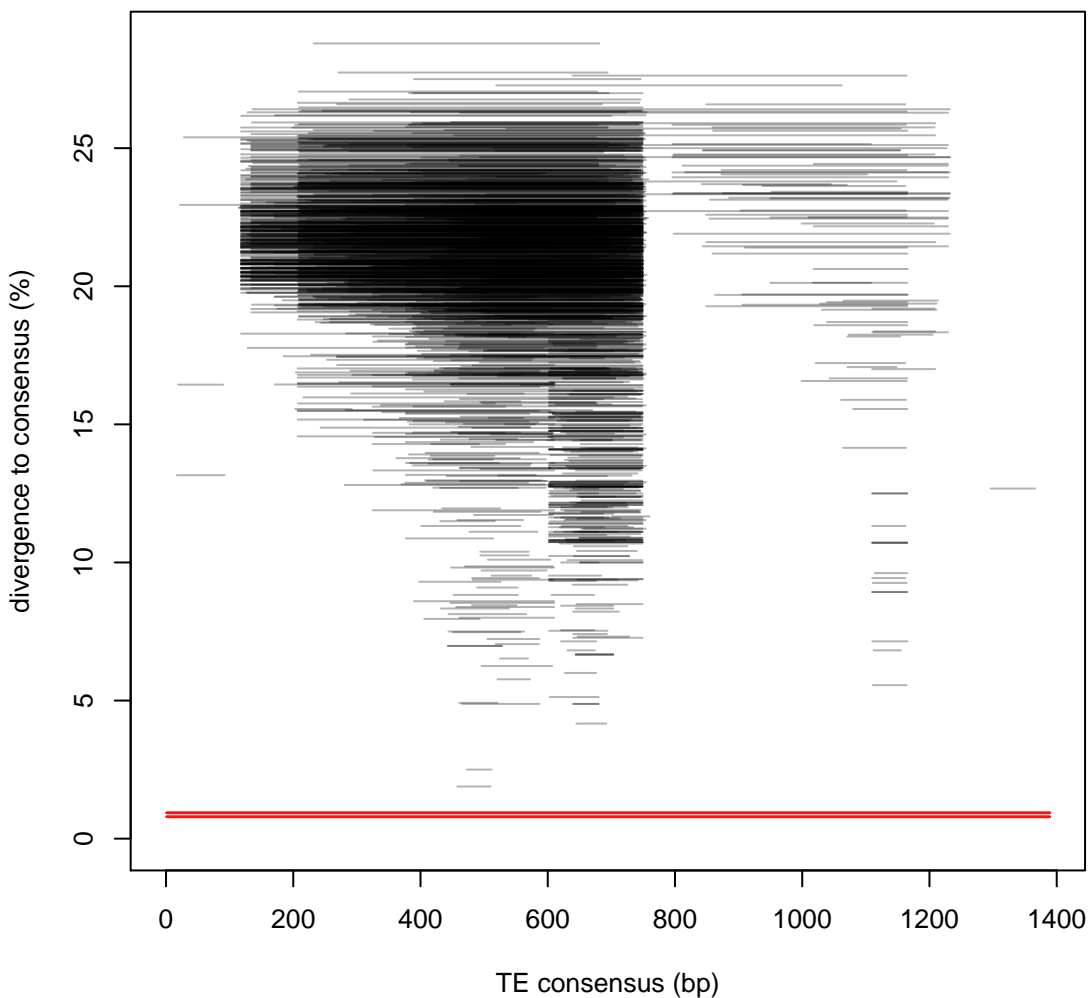
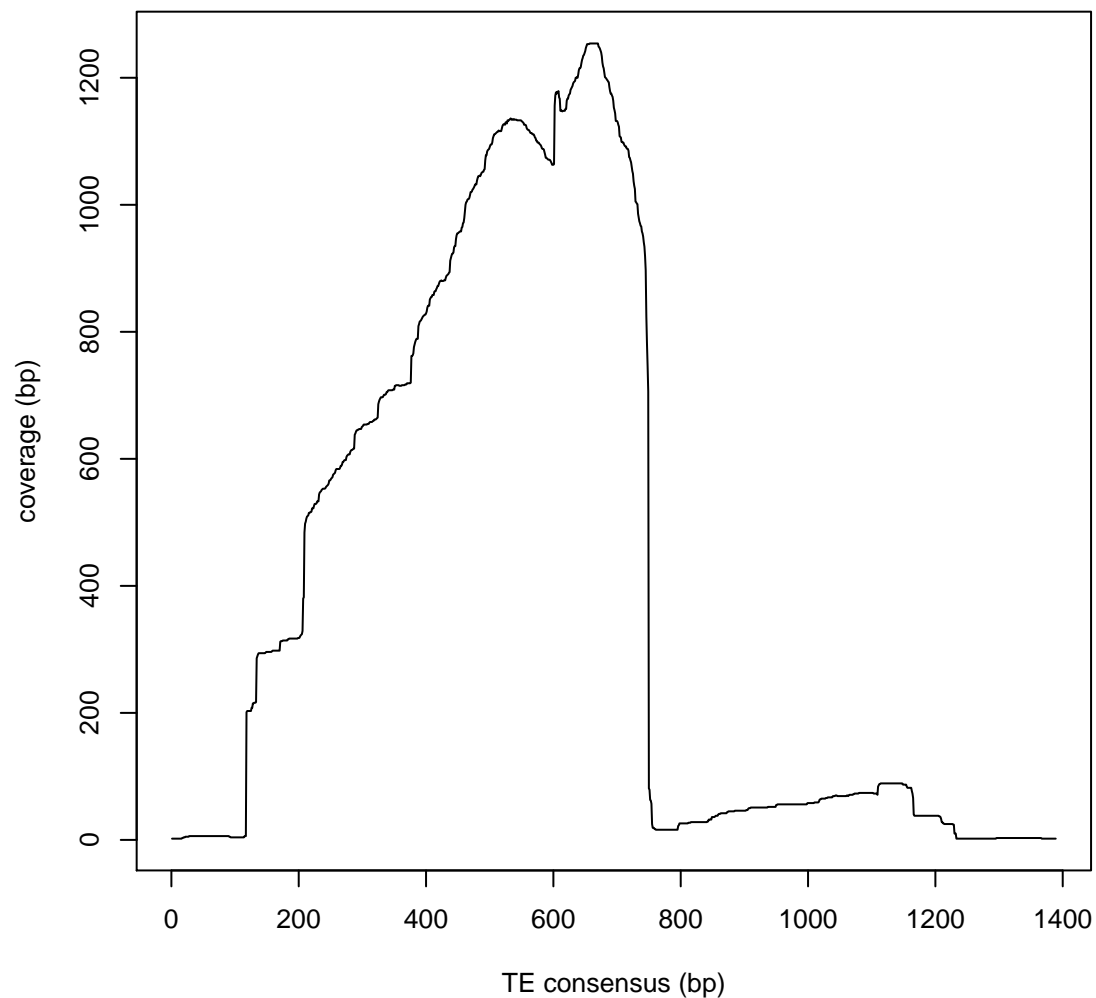


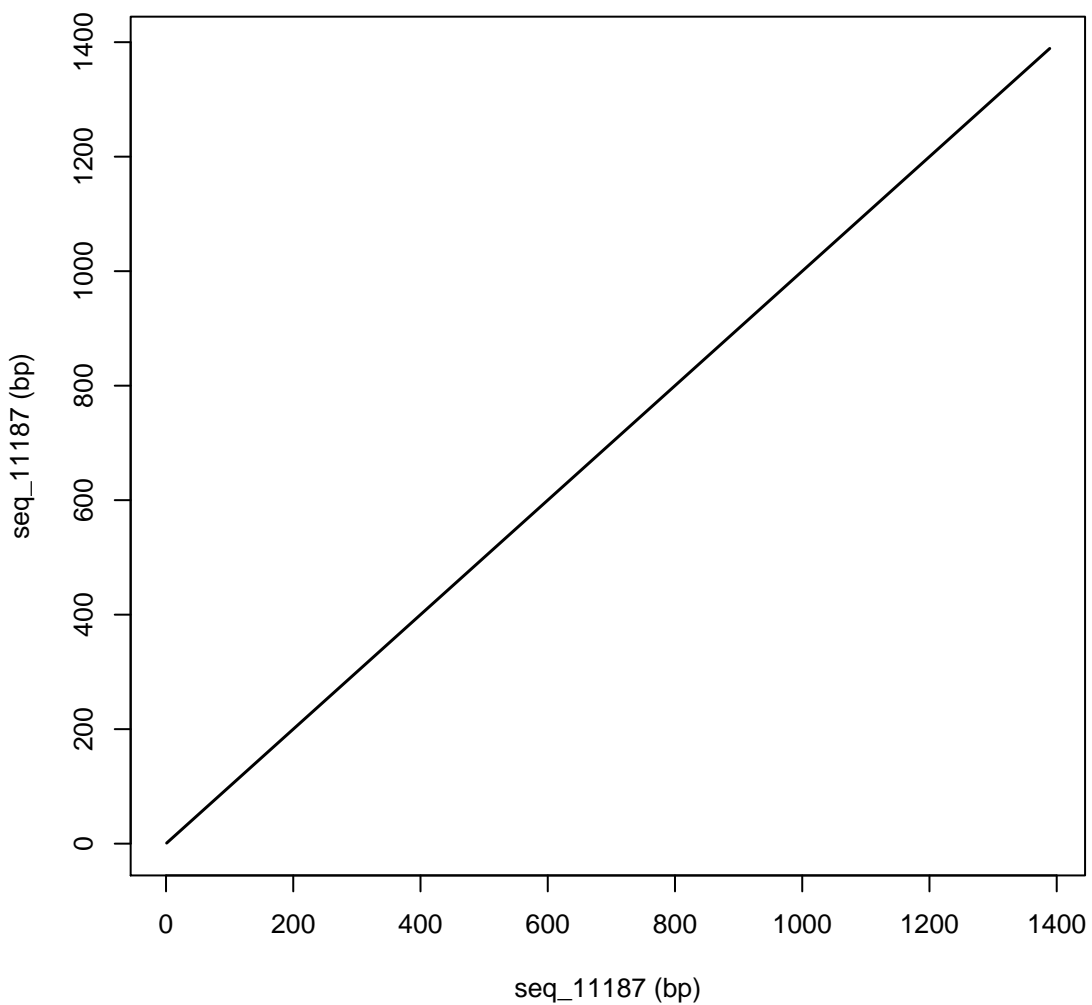
TE: seq\_11187  
consensus size: 1389bp; fragments: 1496; full length: 2 ( $\geq 1250.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

