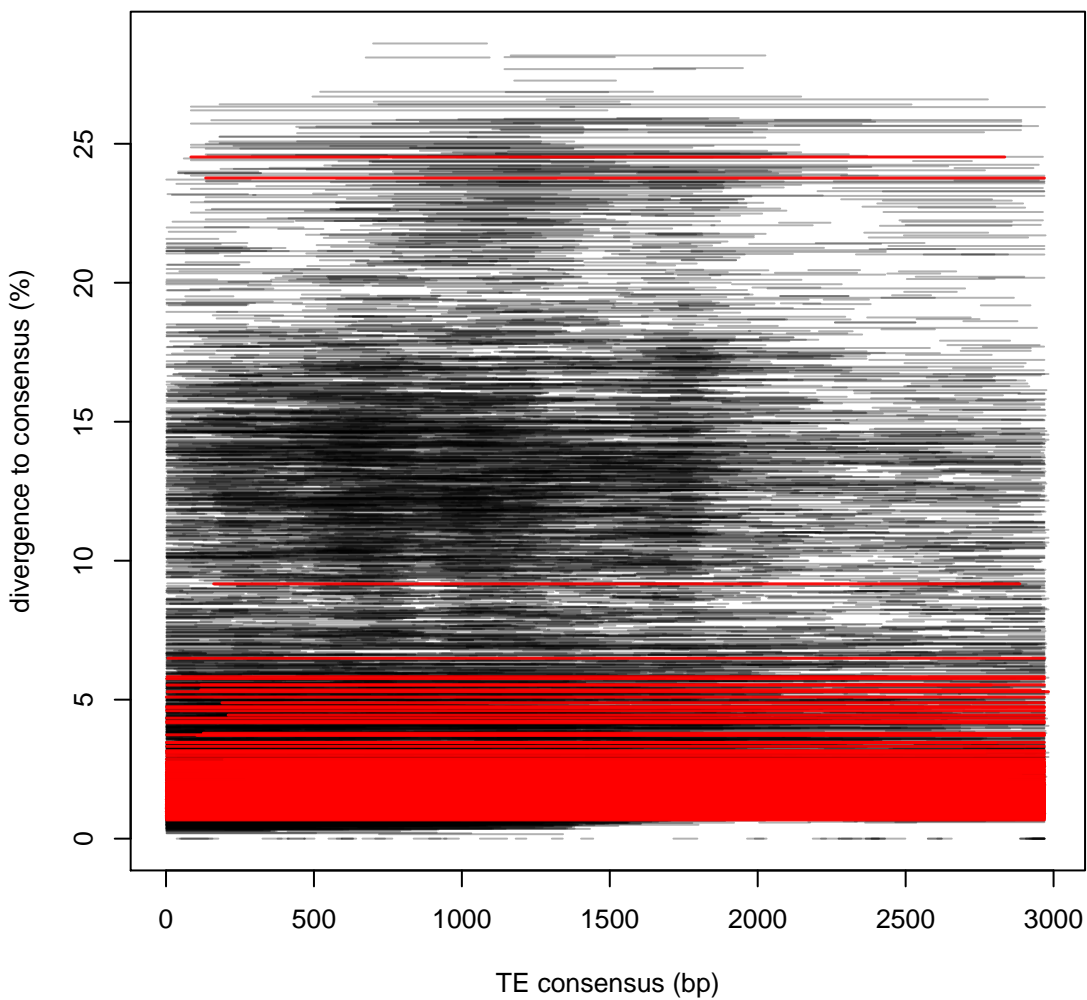
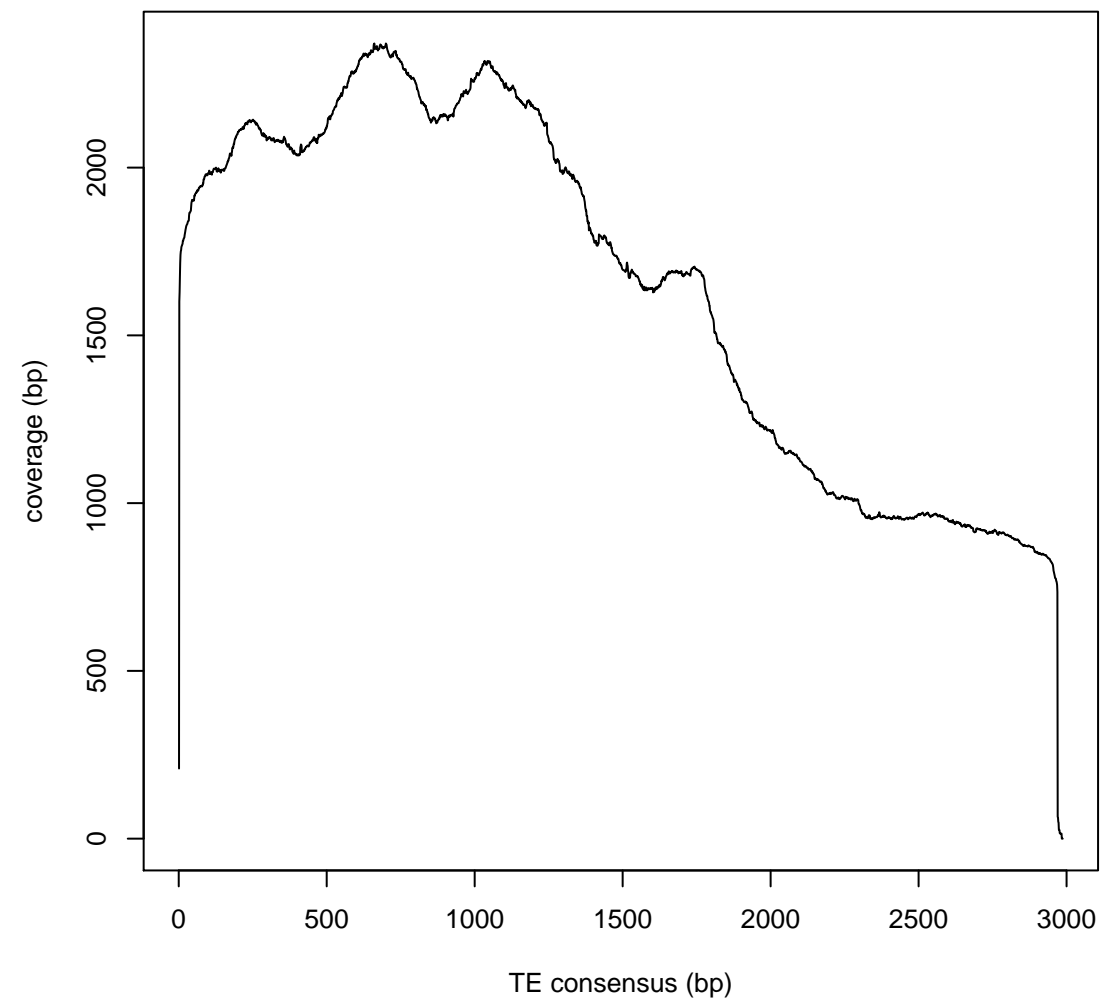


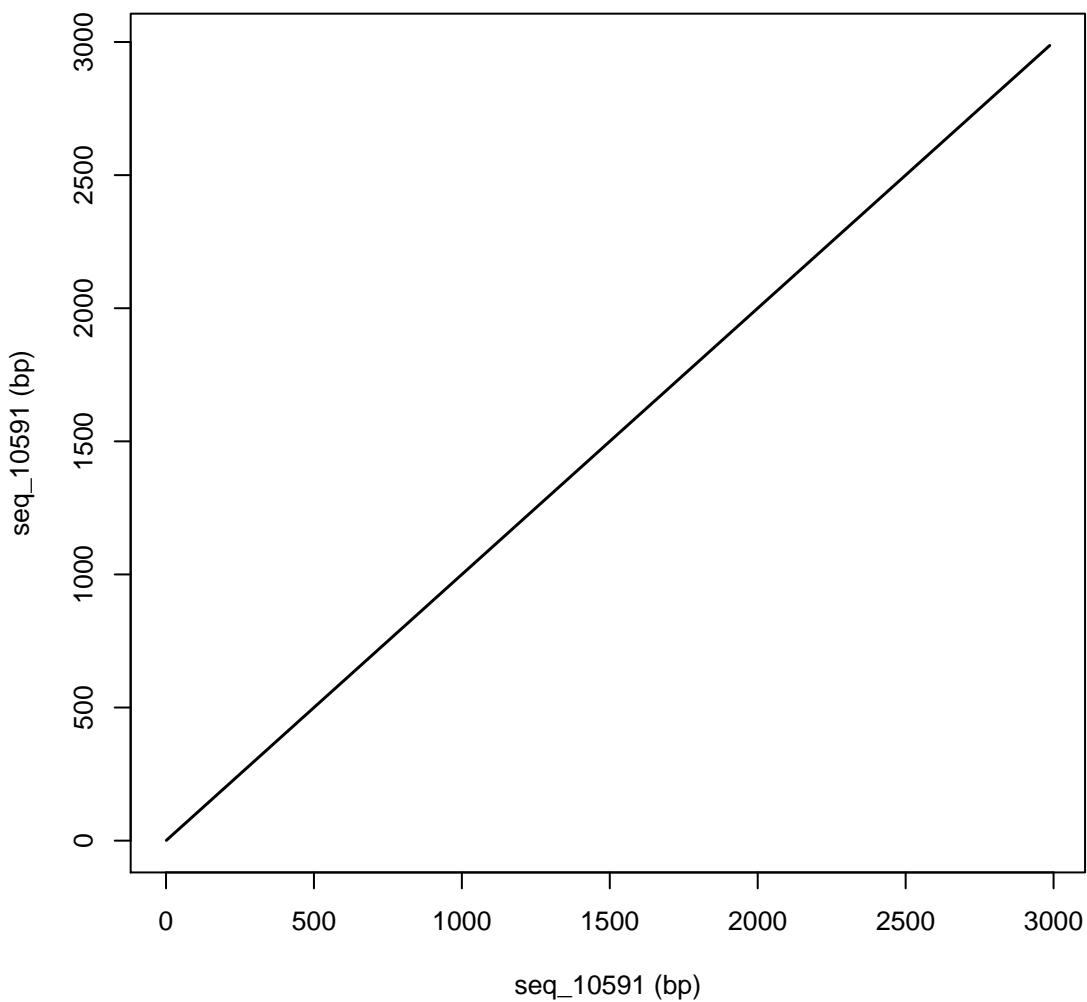
TE: seq\_10591  
consensus size: 2987bp; fragments: 7606; full length: 280 ( $\geq 2688.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

