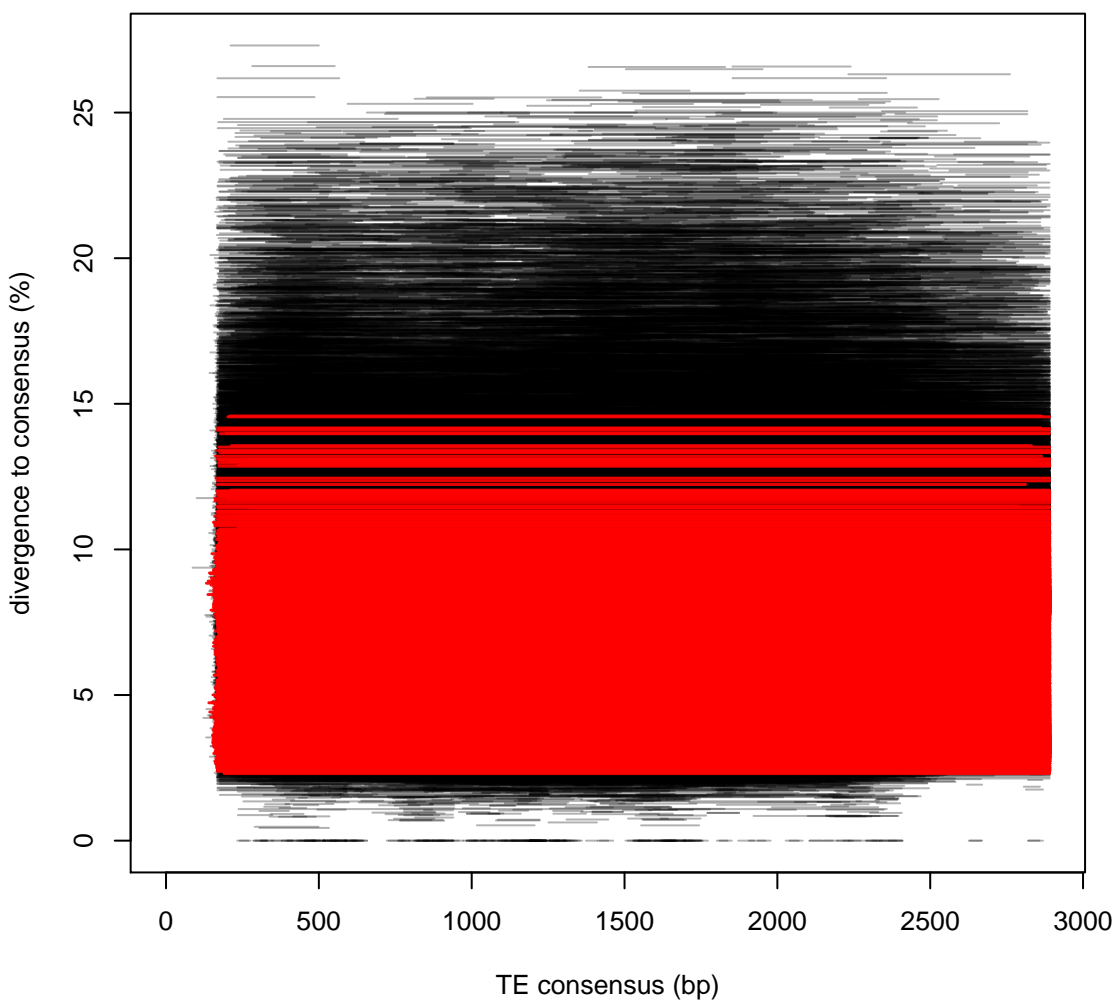
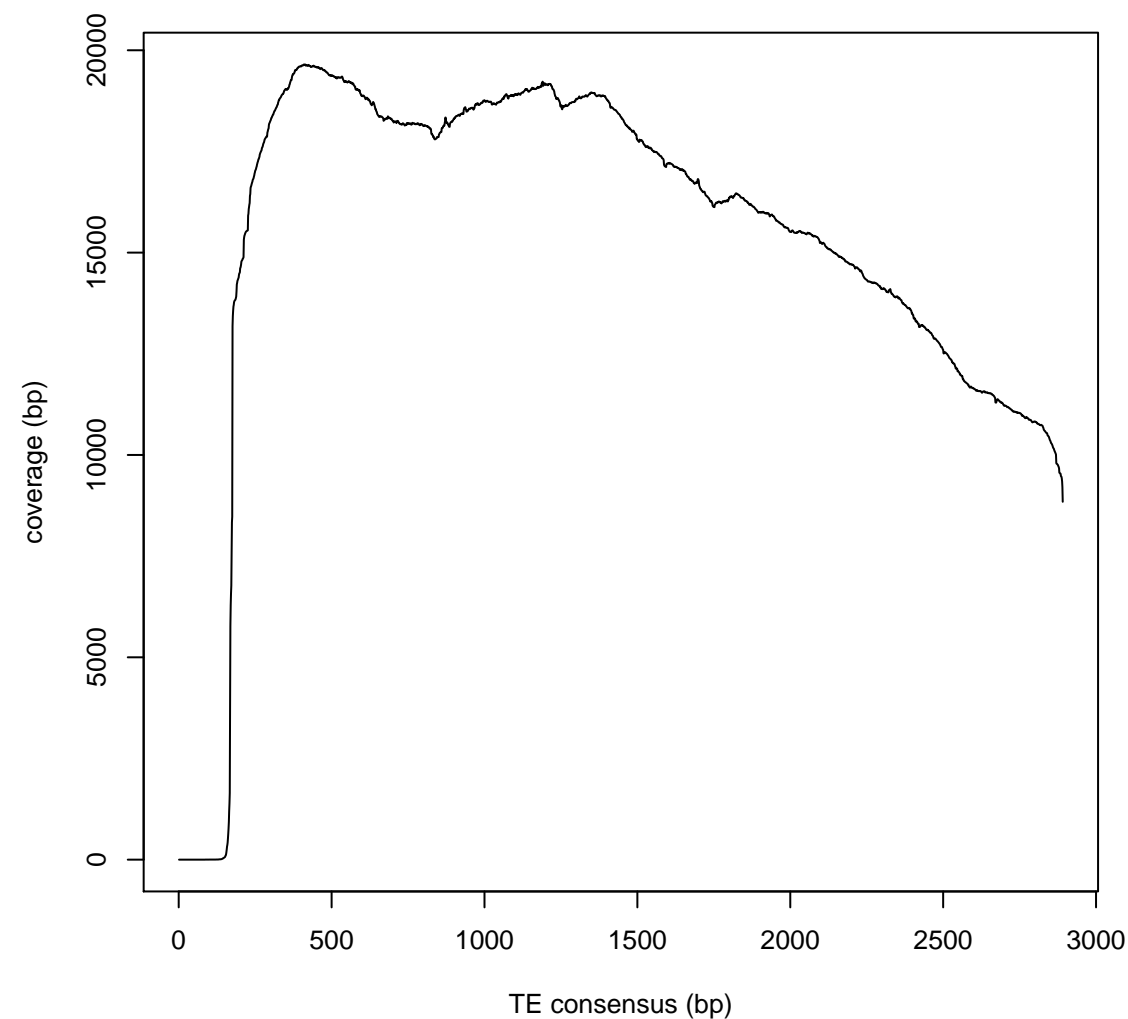


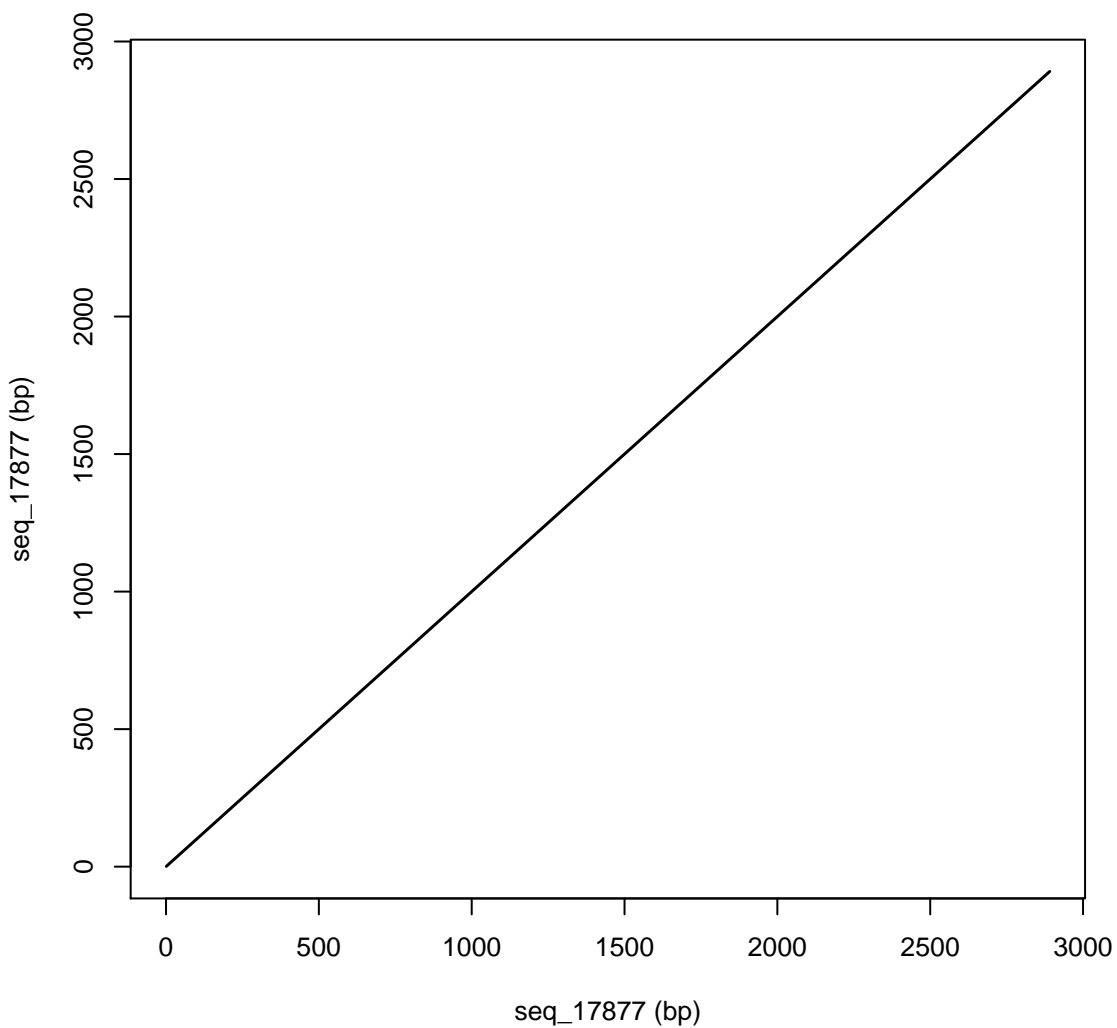
TE: seq\_17877  
consensus size: 2891bp; fragments: 54500; full length: 3086 ( $\geq 2601.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

