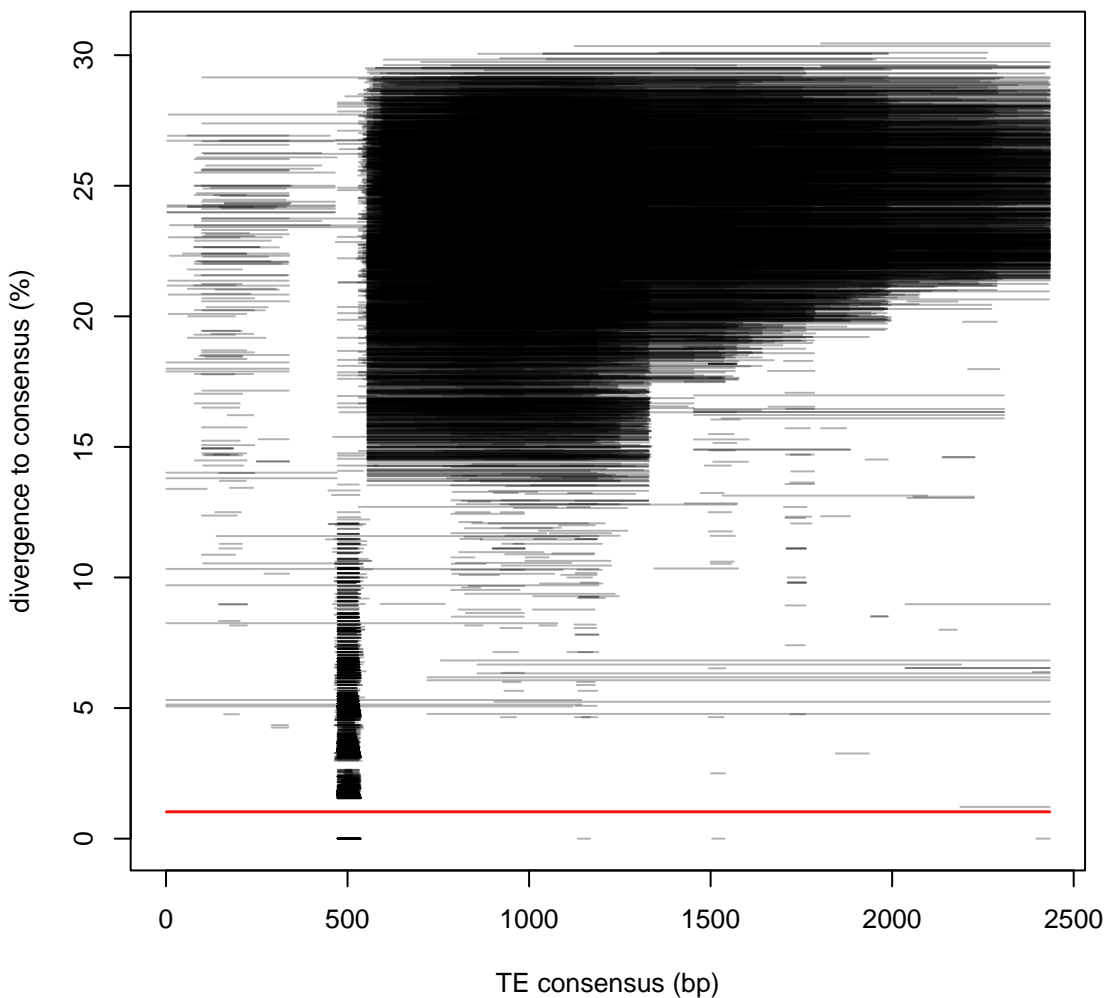
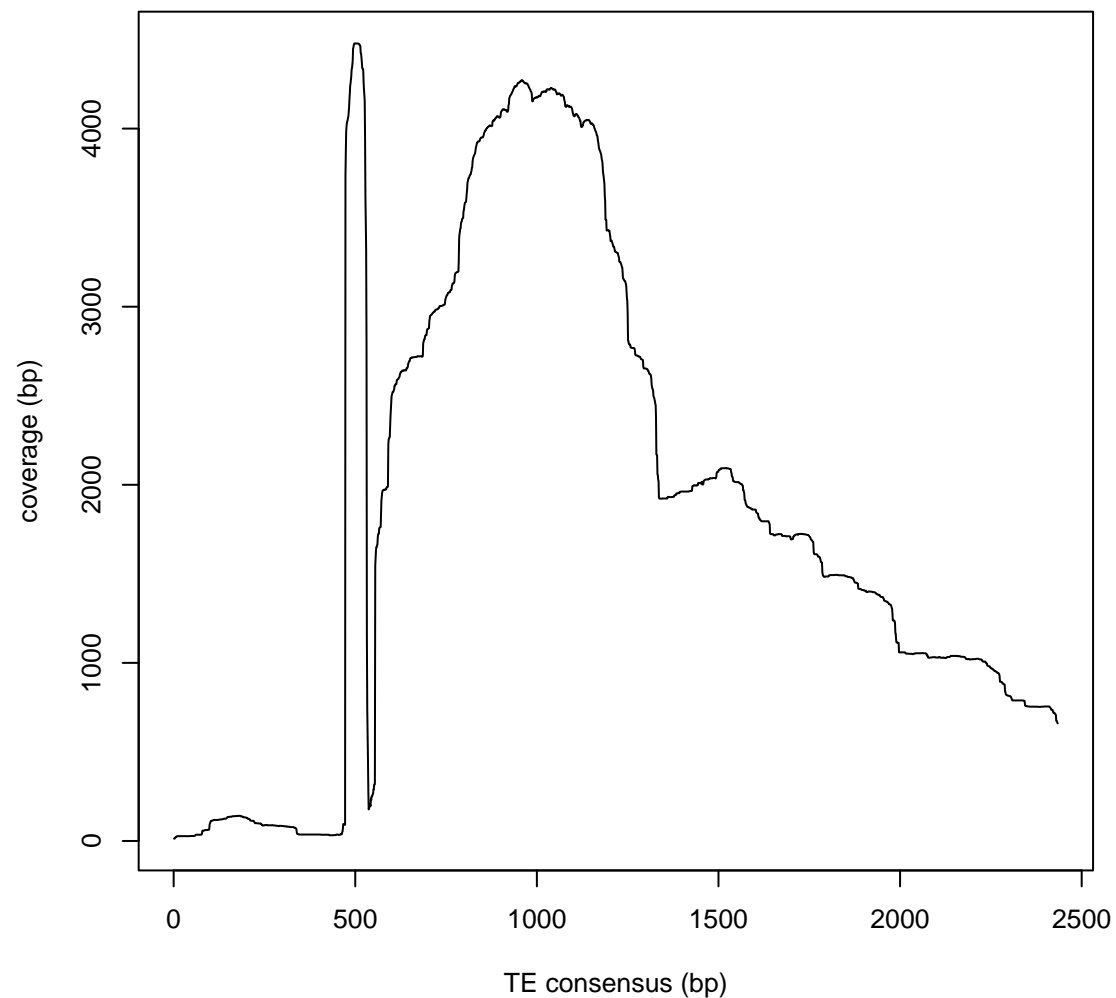


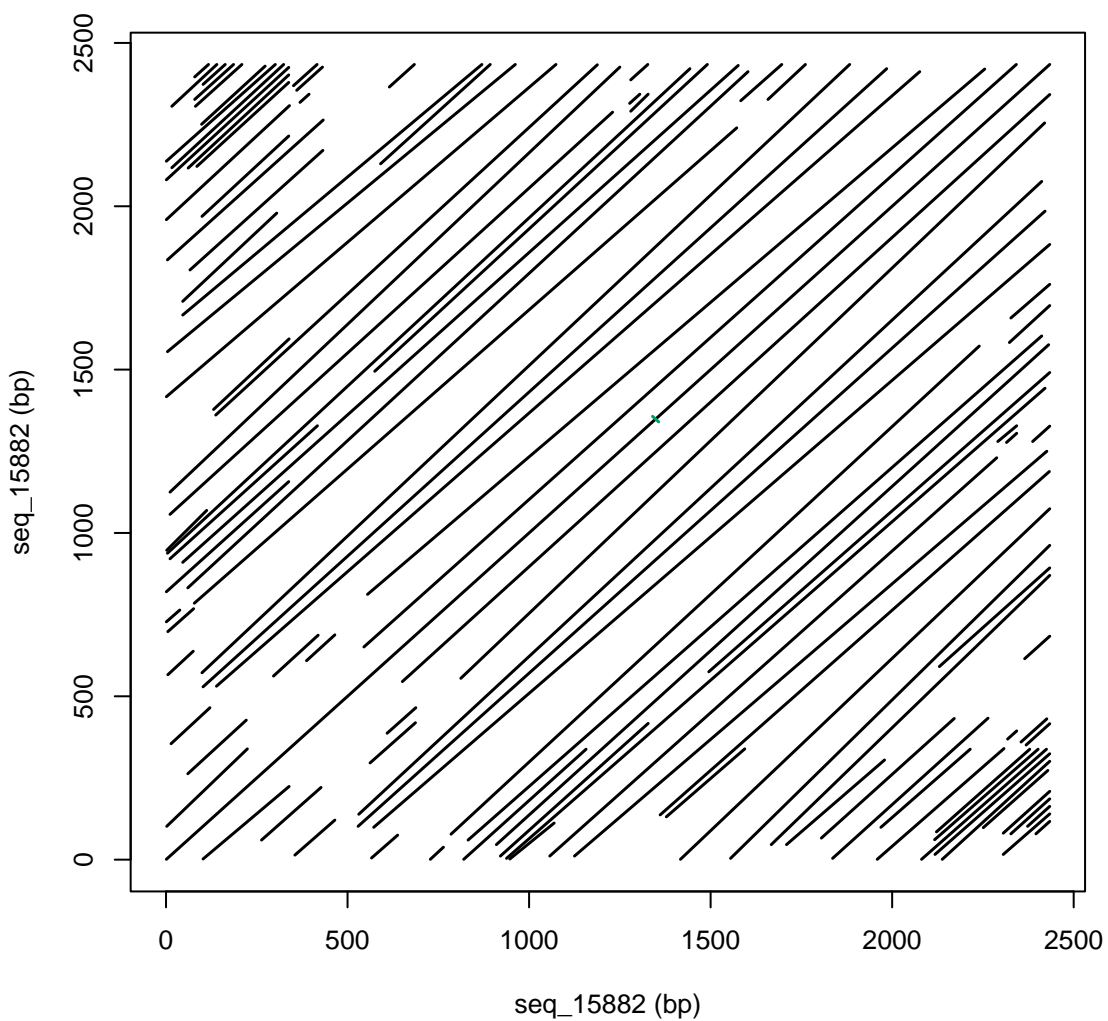
TE: seq\_15882  
consensus size: 2434bp; fragments: 9553; full length: 1 (>=2190.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

