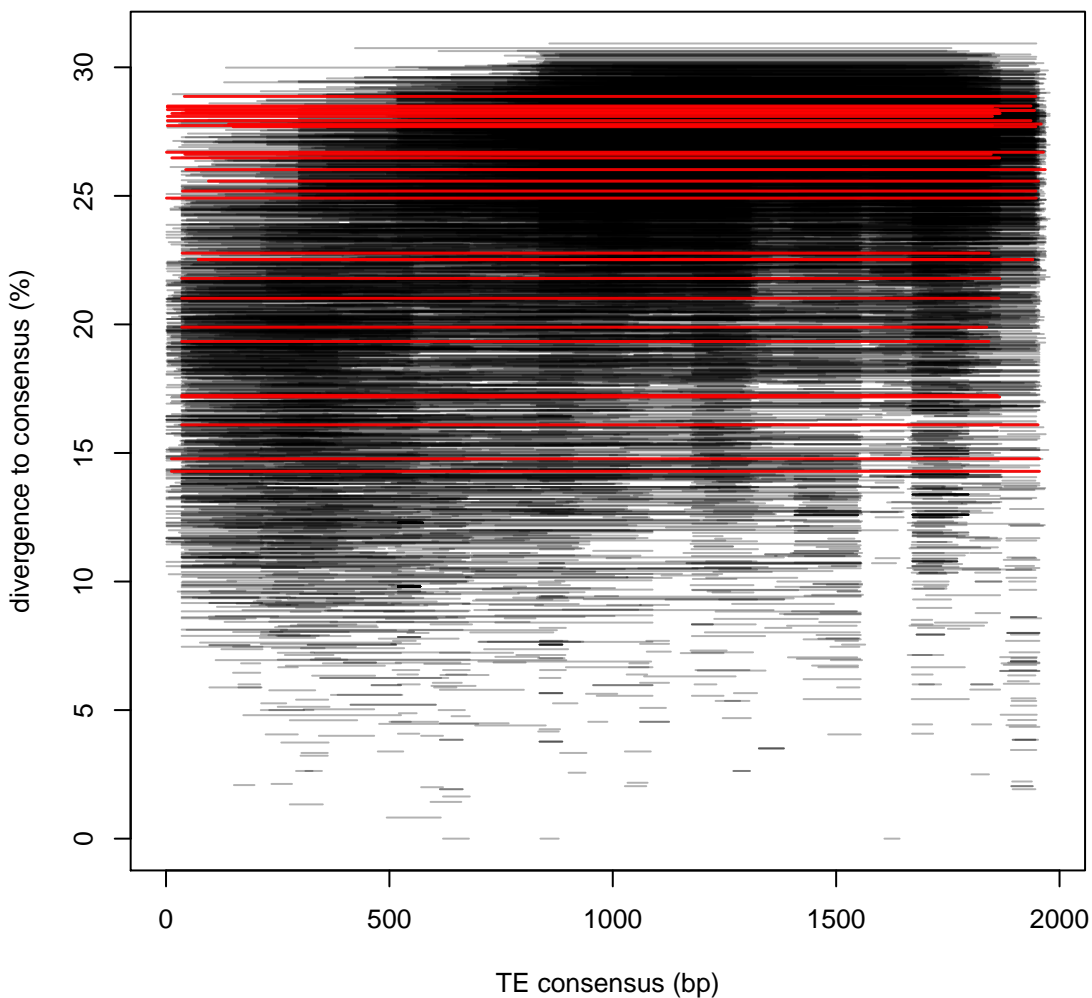
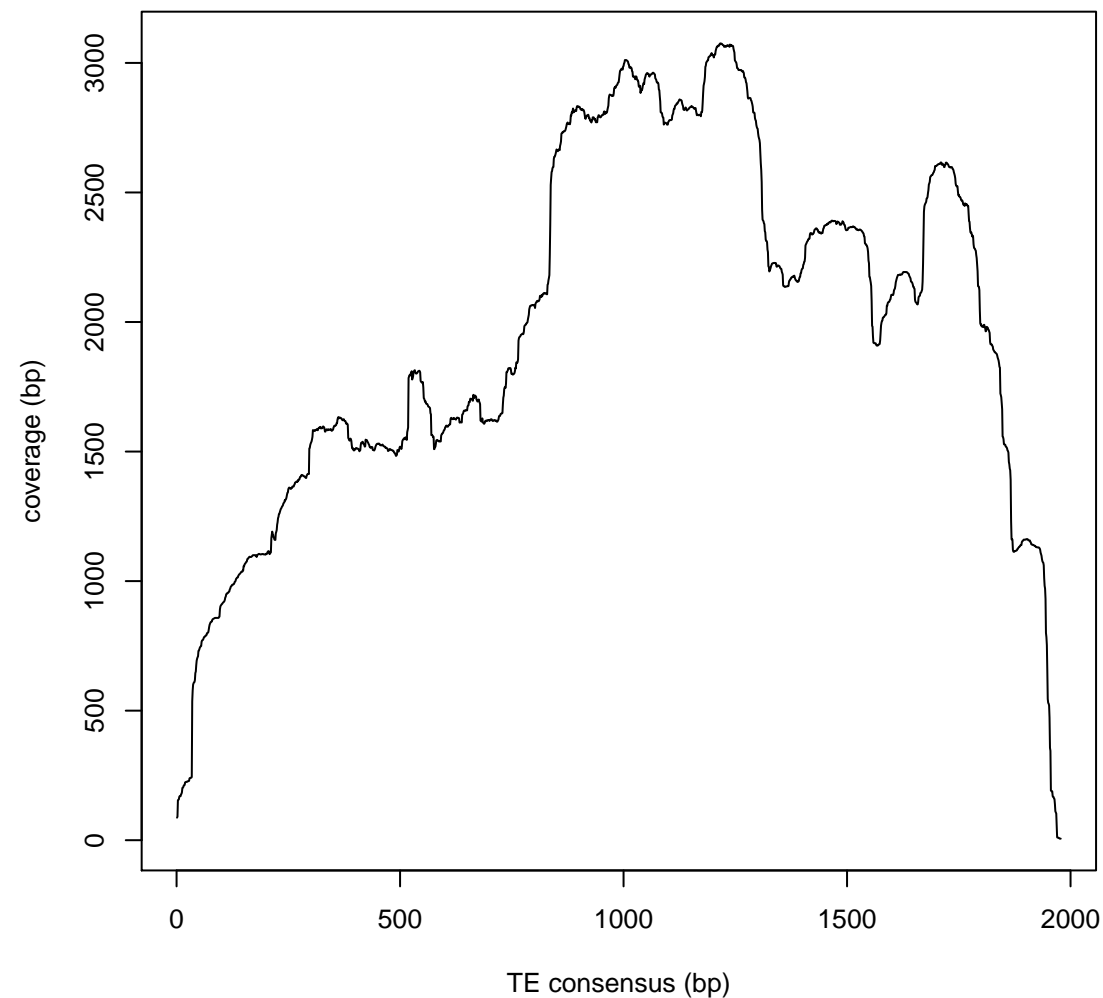


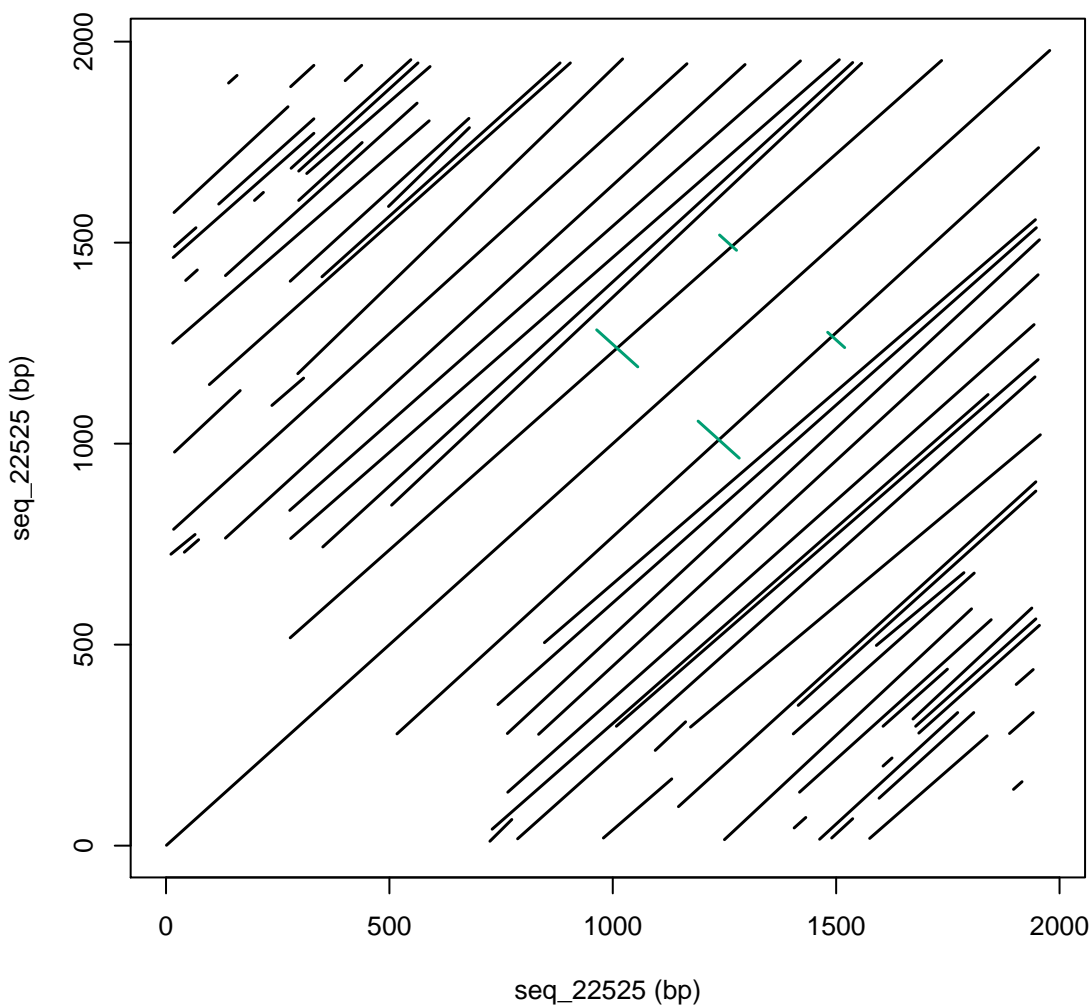
TE: seq\_22525  
consensus size: 1978bp; fragments: 7331; full length: 31 (>=1780.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

