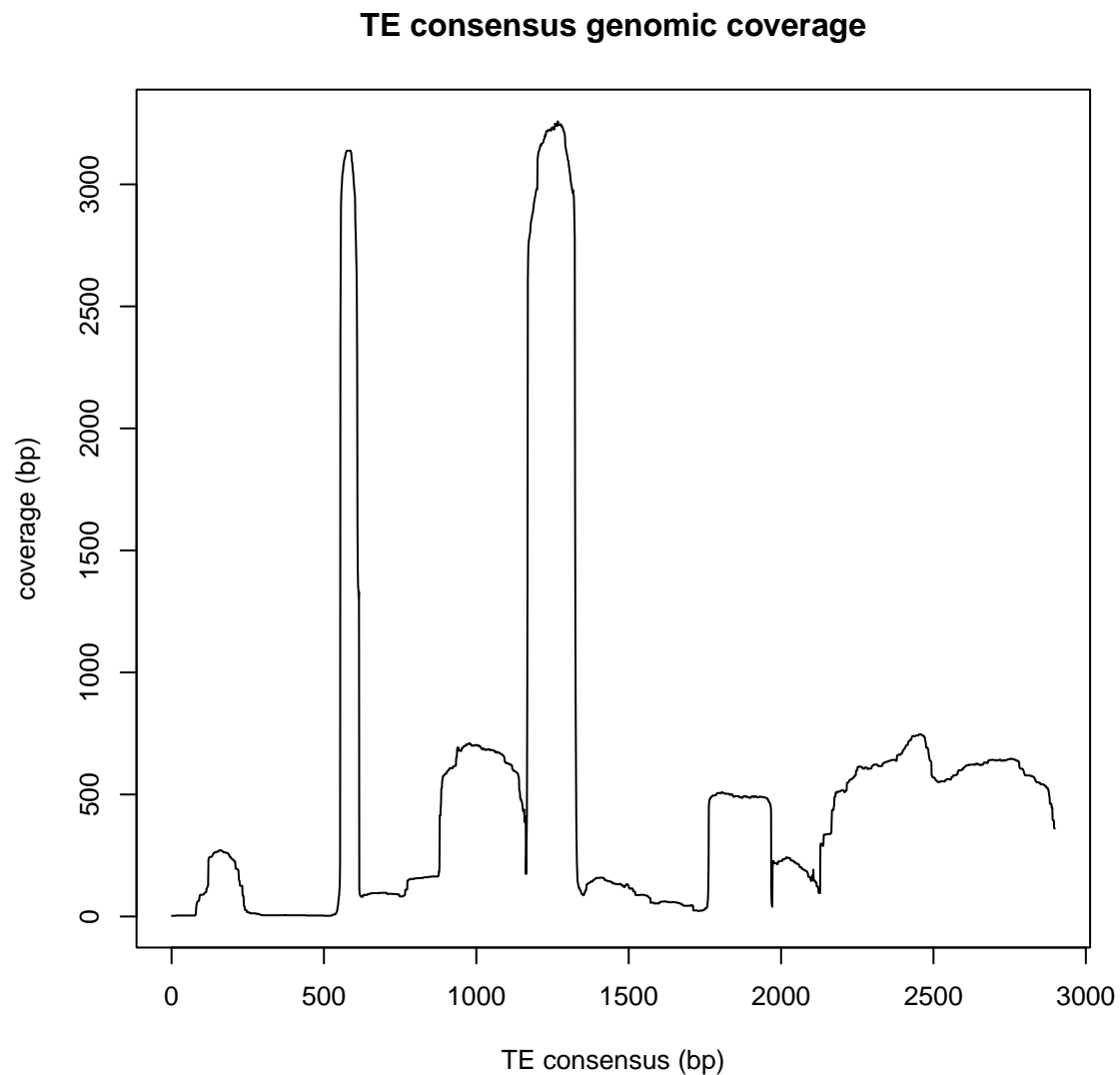
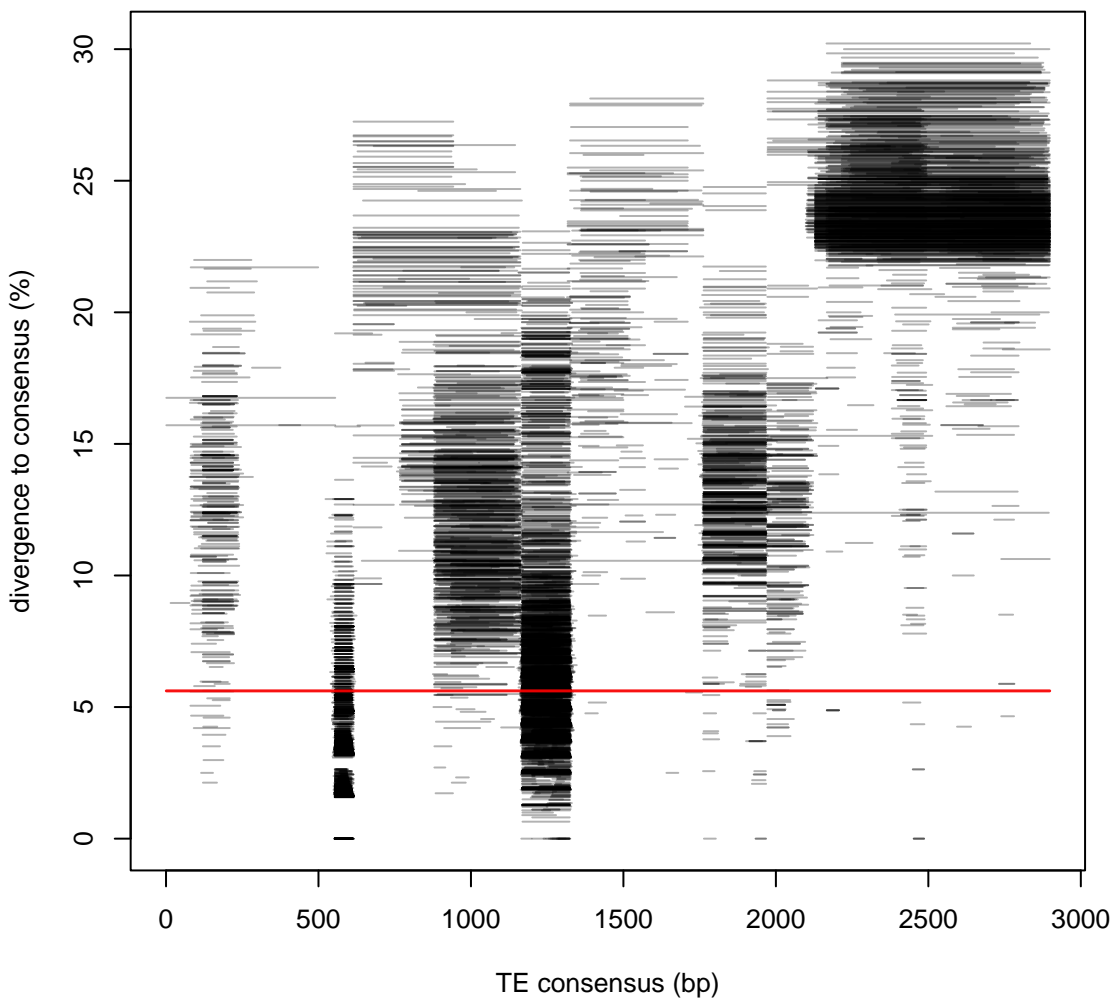
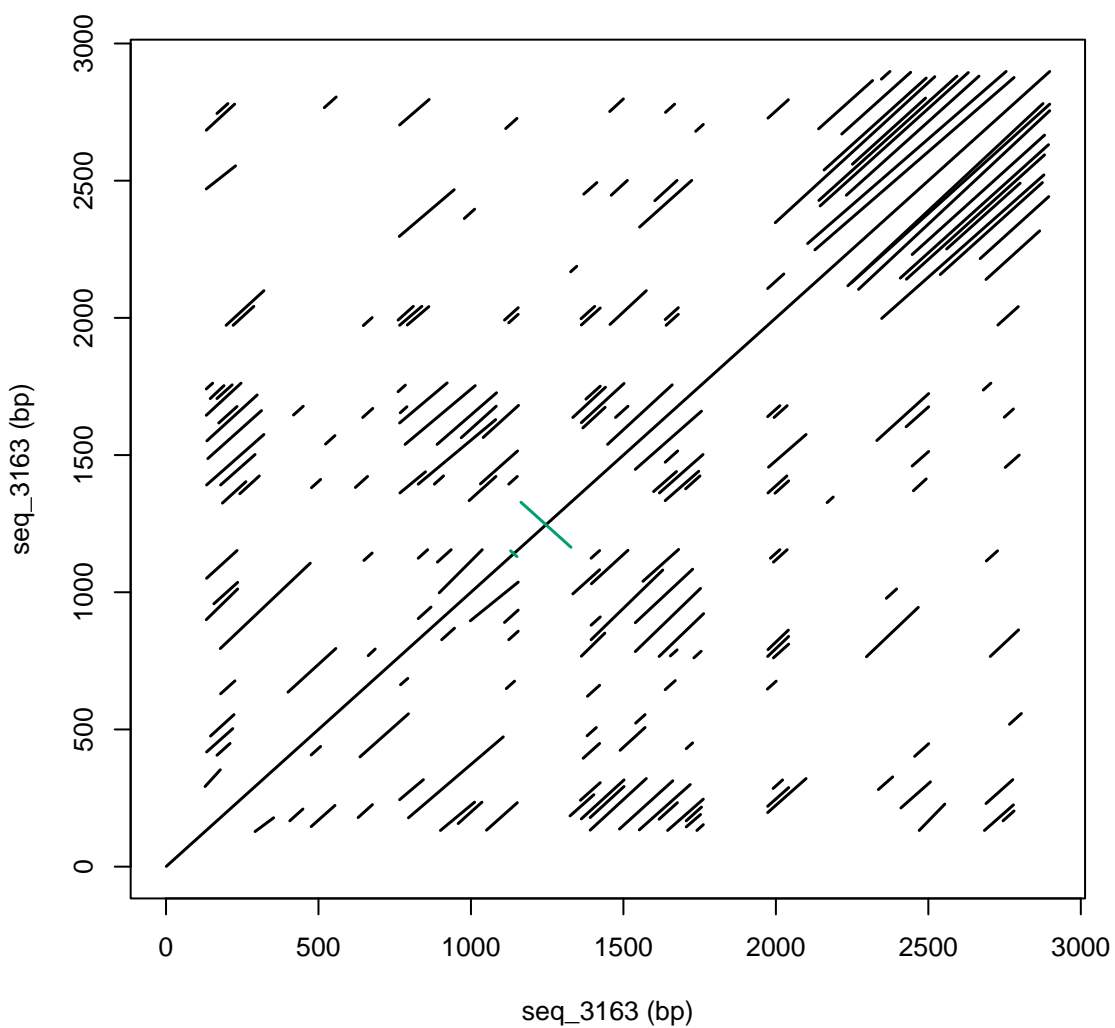


TE: seq\_3163  
consensus size: 2898bp; fragments: 9570; full length: 1 (>=2608.2bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

