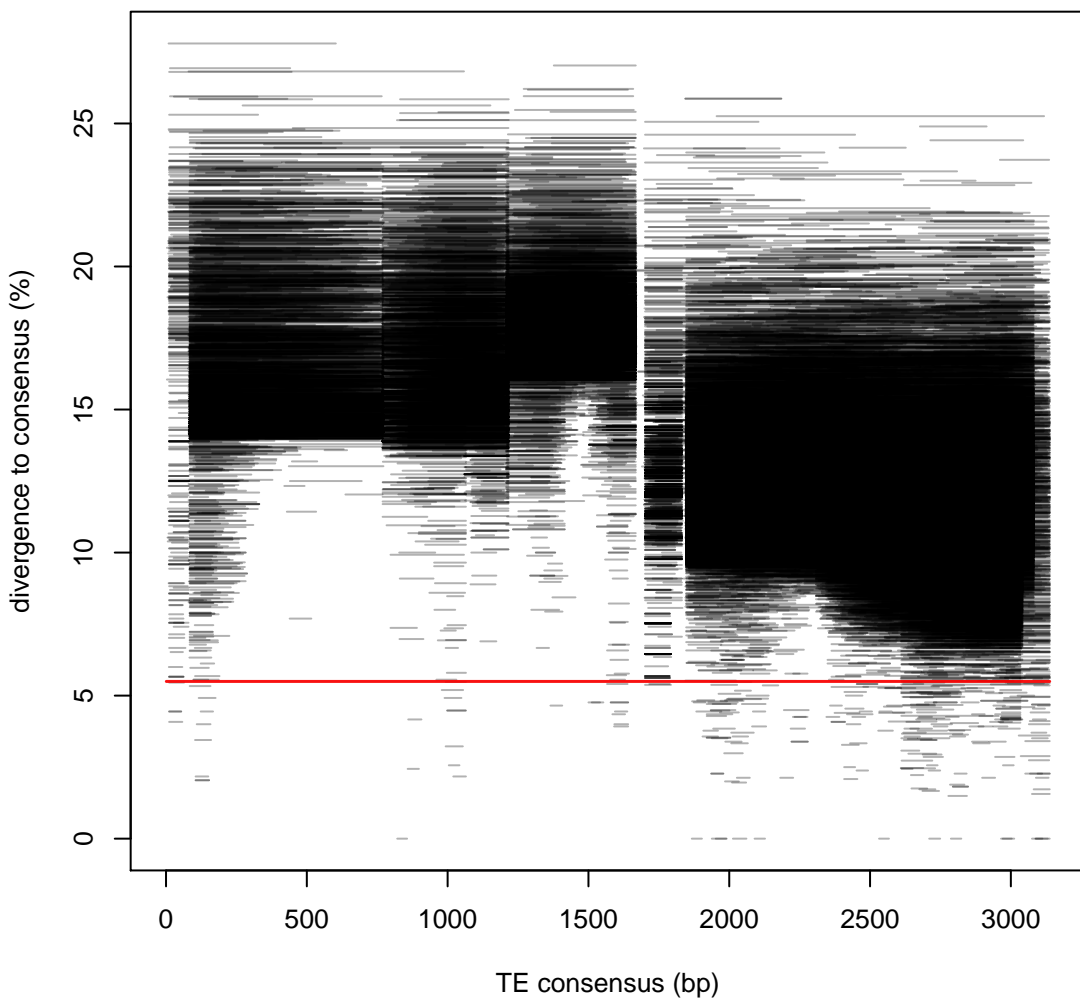
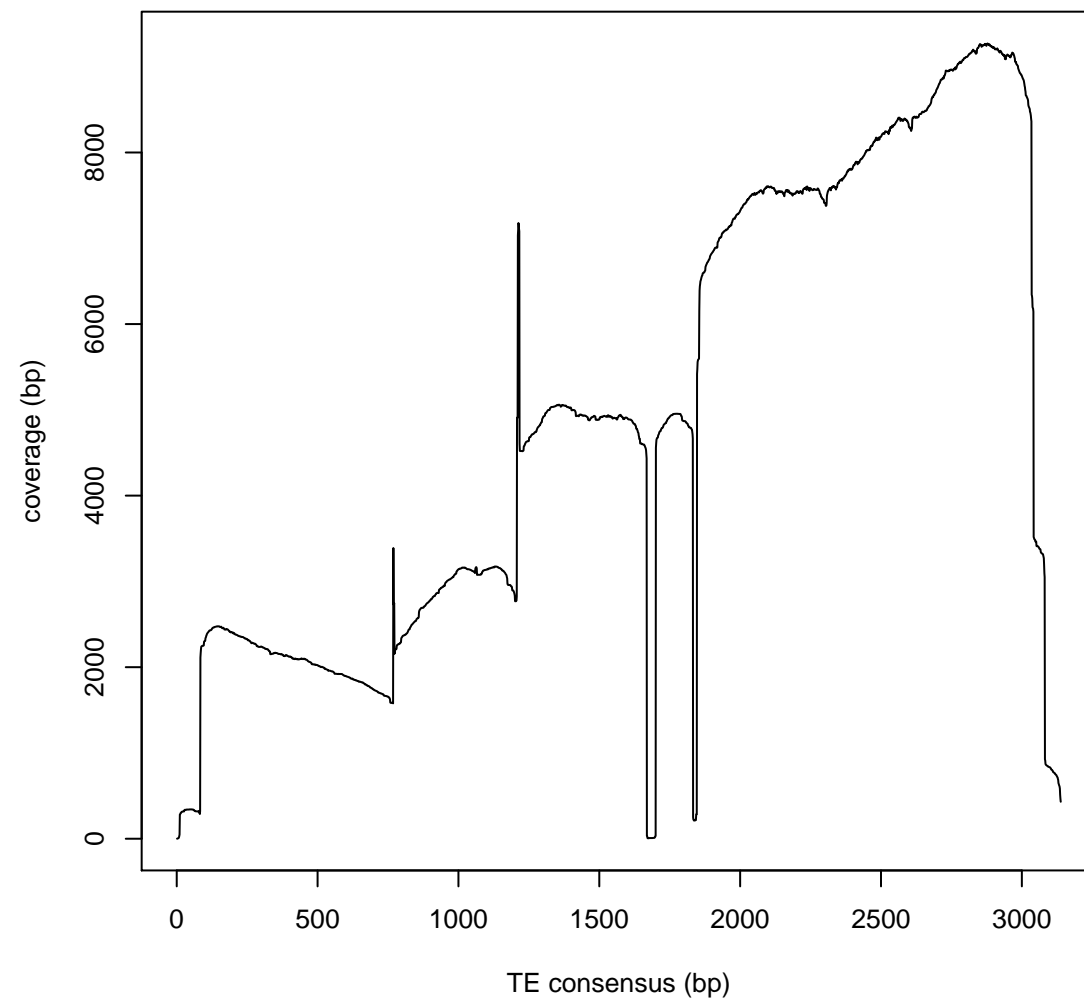


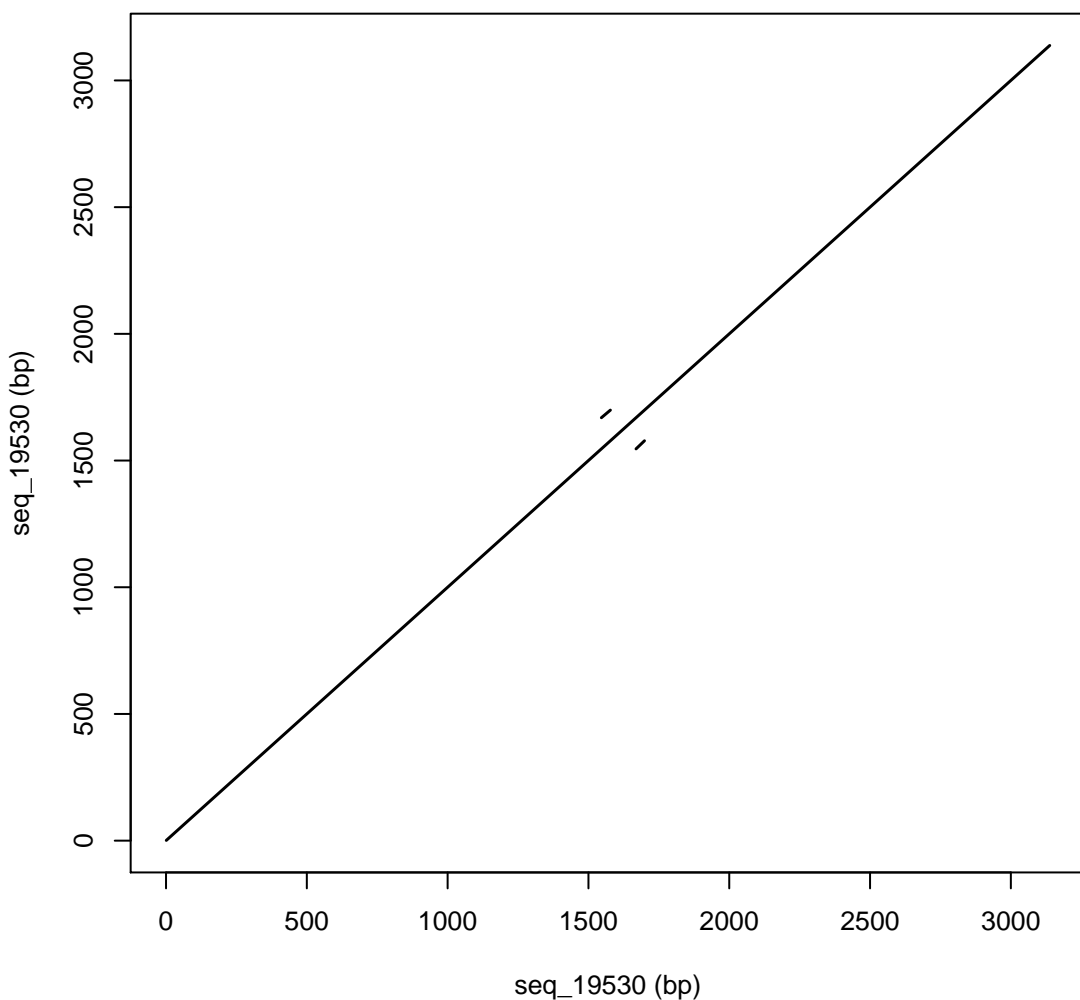
TE: seq\_19530  
consensus size: 3138bp; fragments: 31736; full length: 1 (>=2824.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

