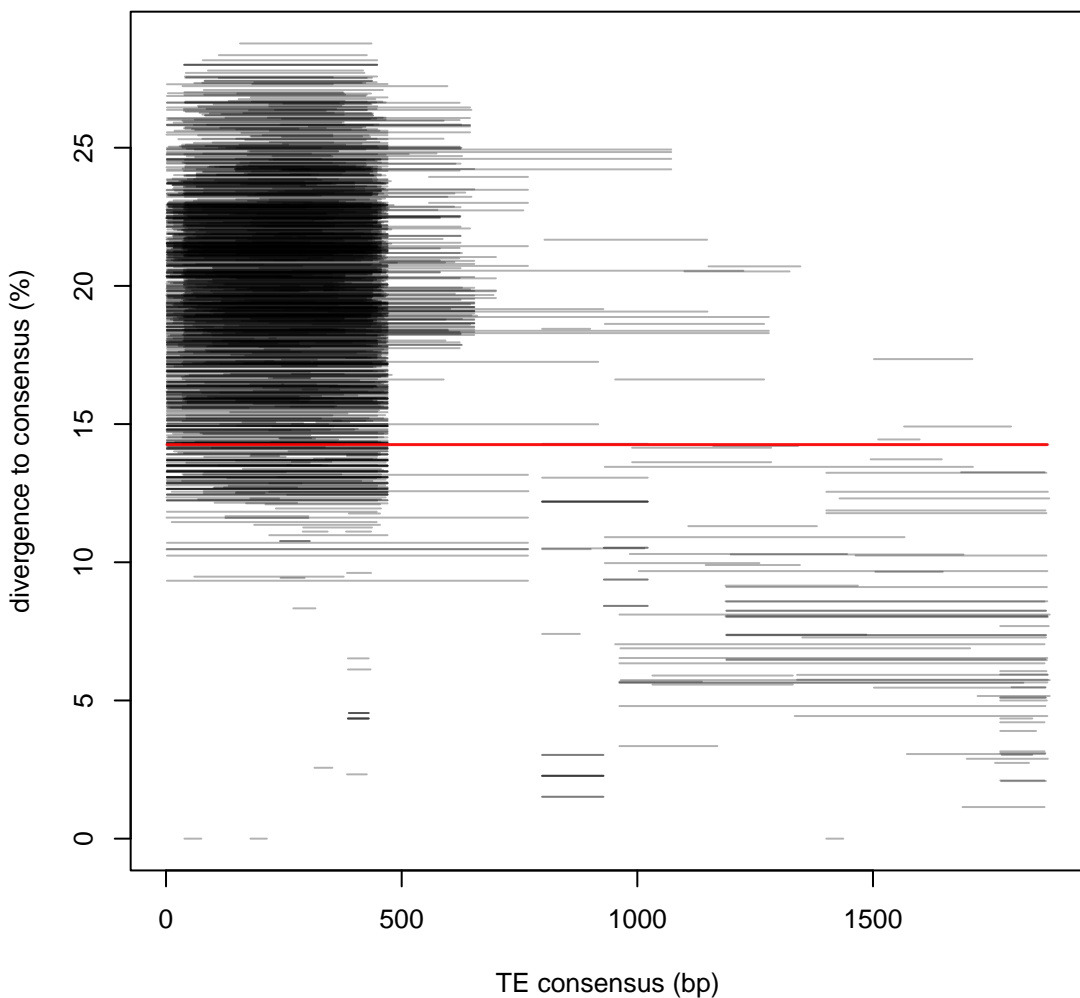
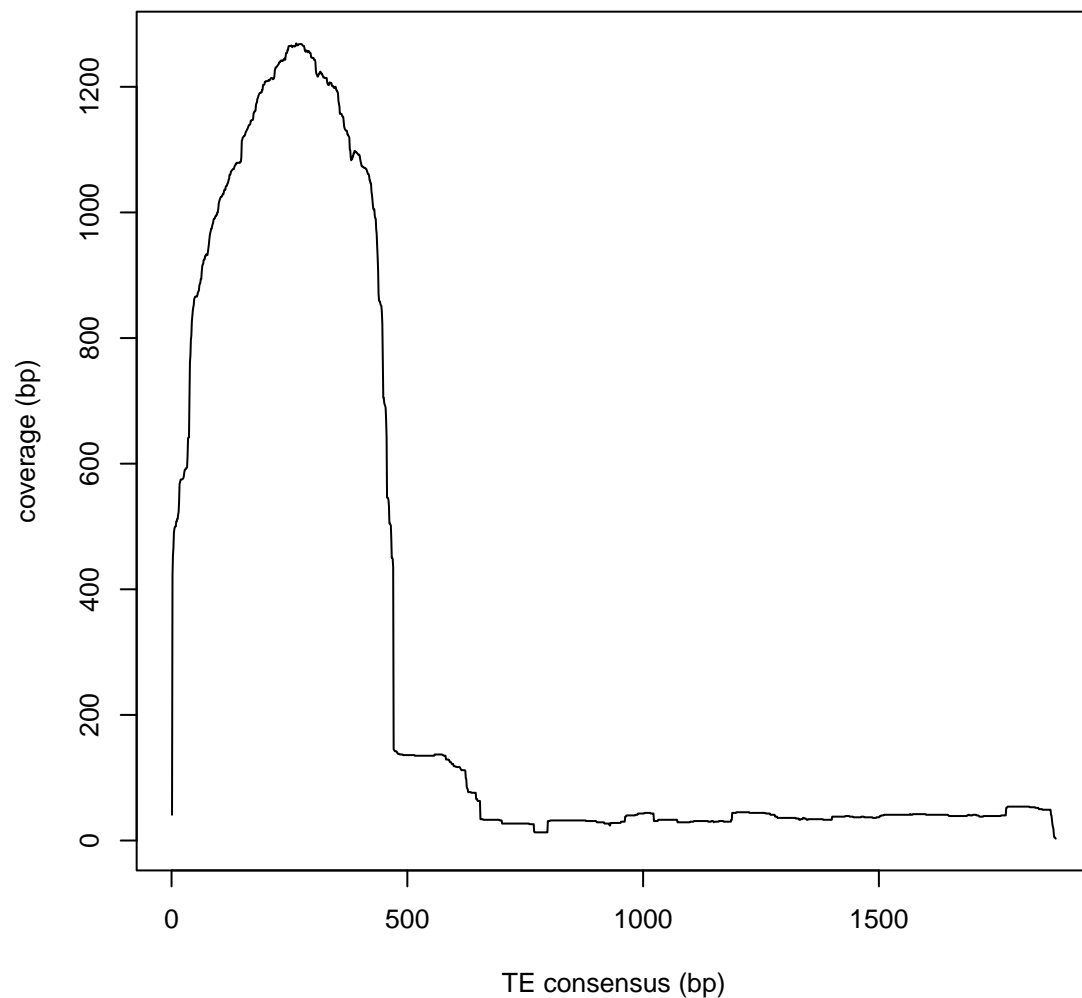


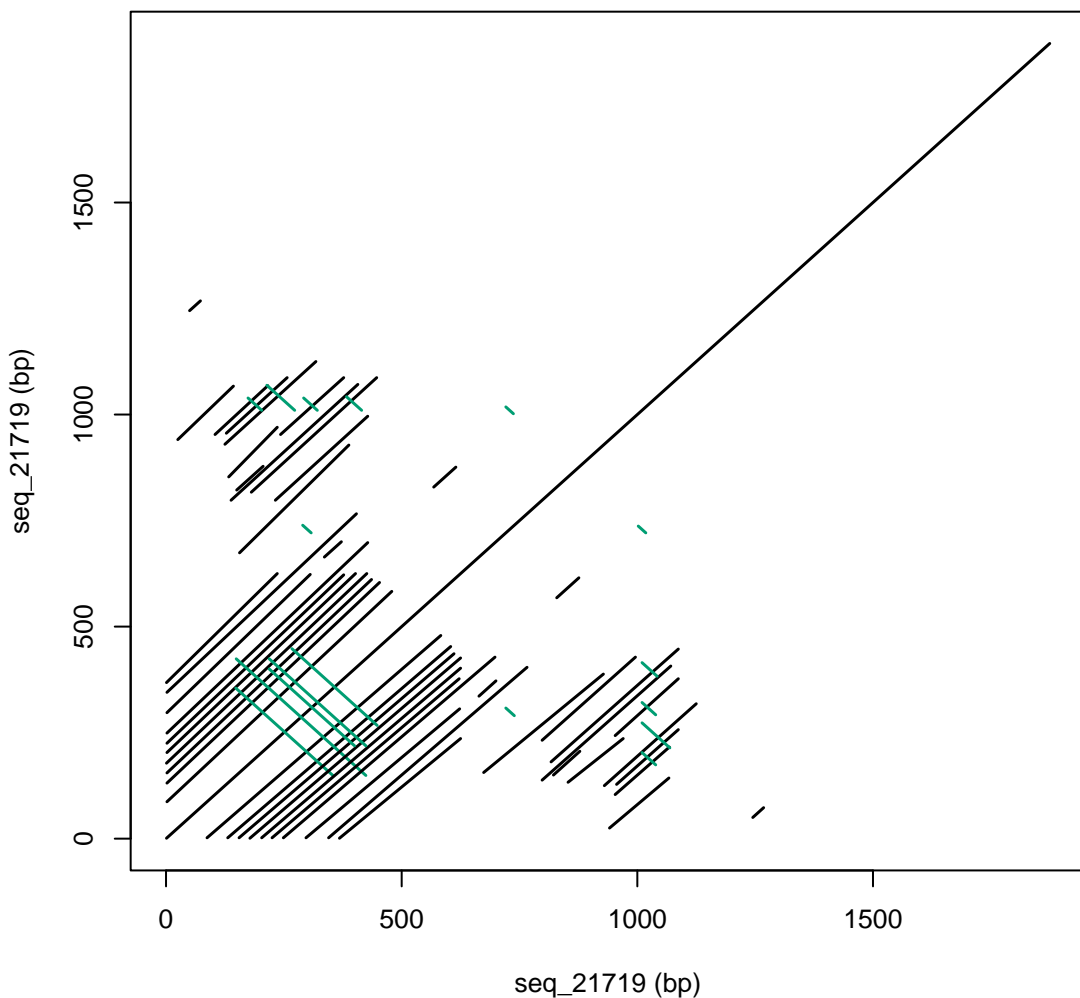
TE: seq\_21719  
consensus size: 1875bp; fragments: 1448; full length: 1 (>=1687.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

