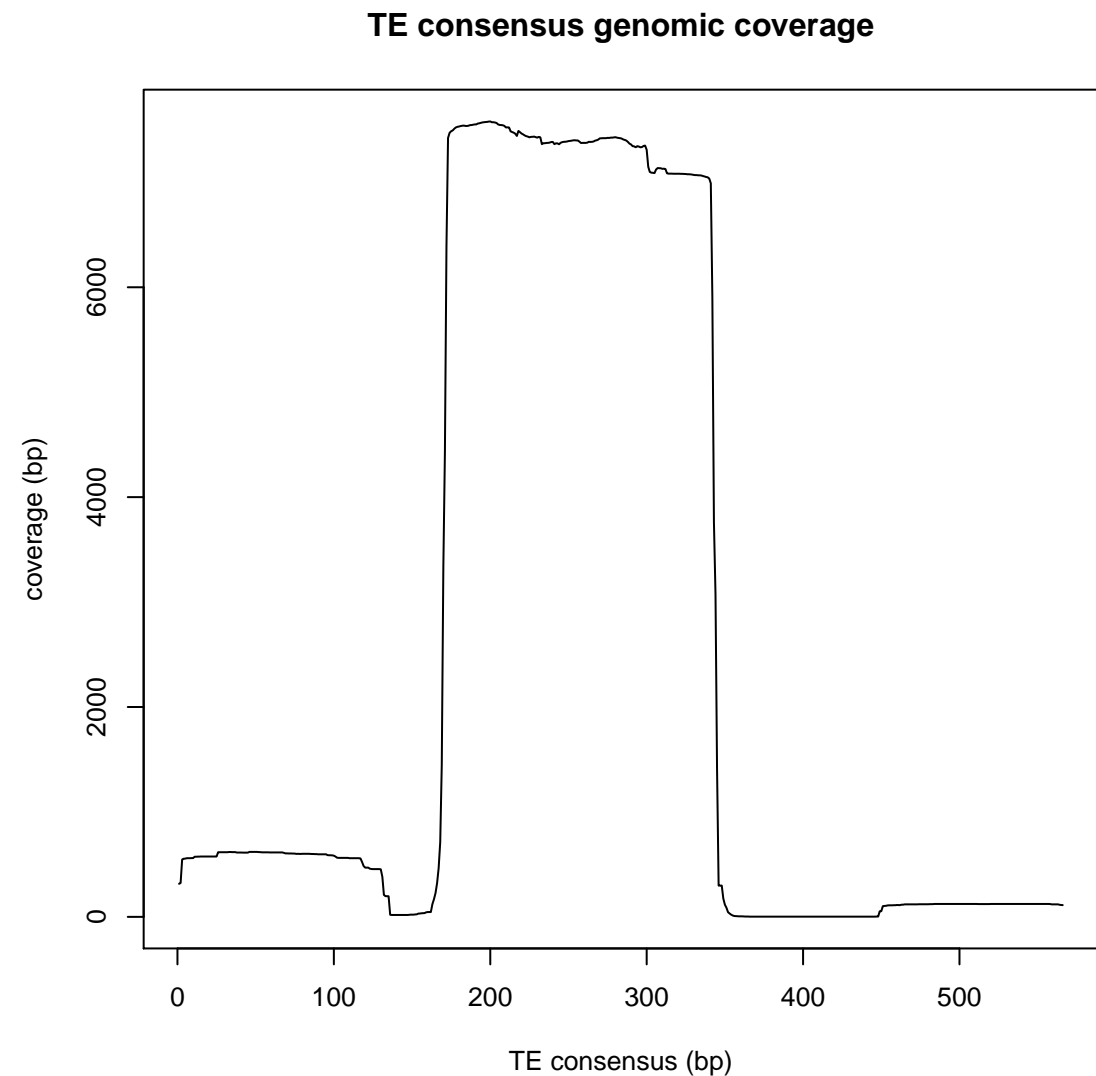
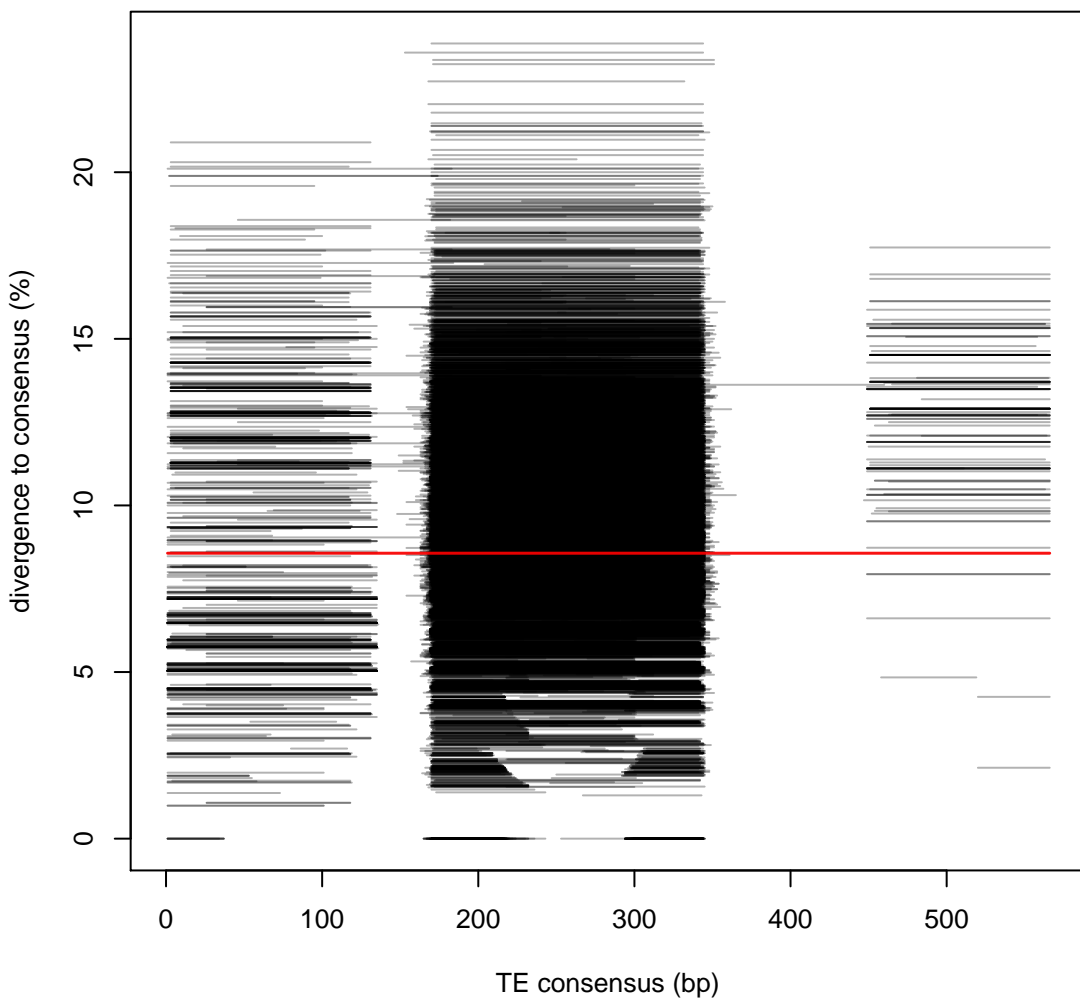
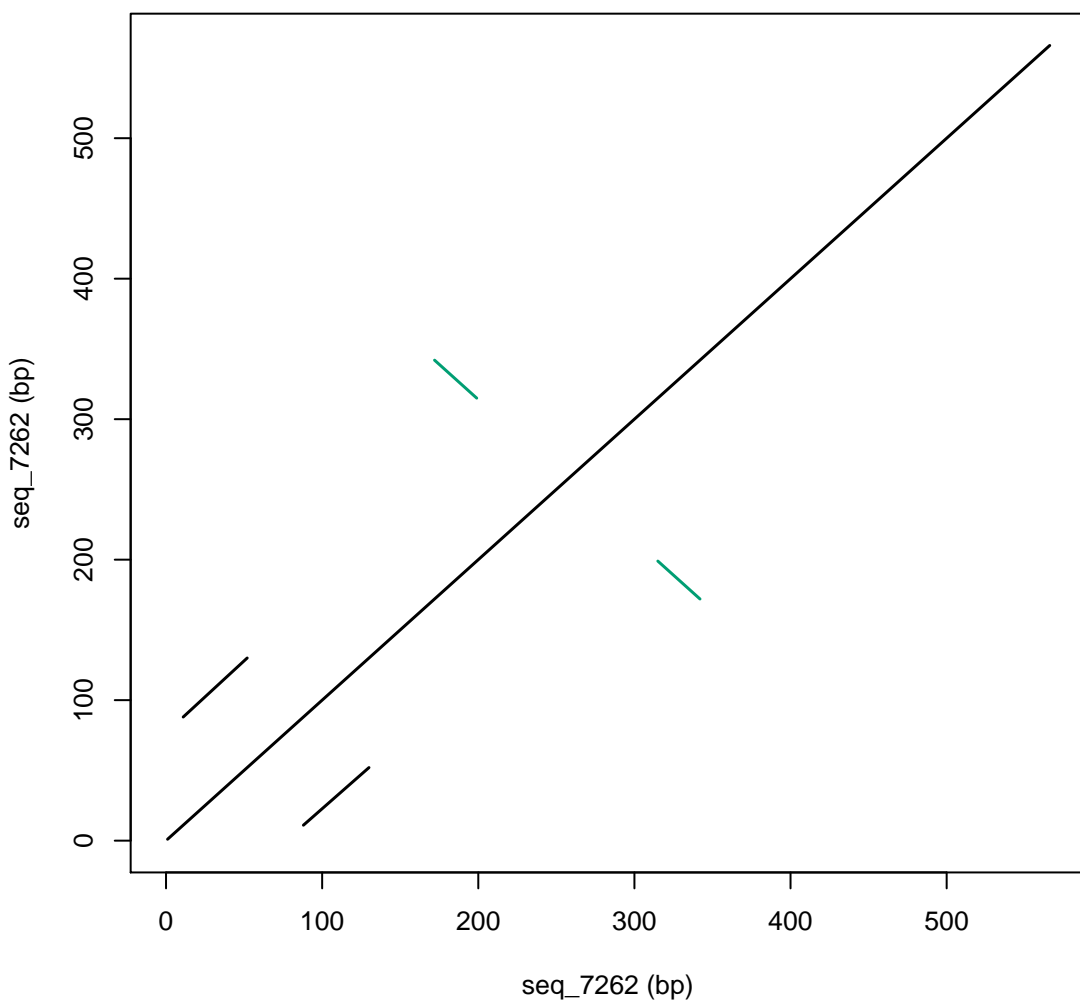


TE: seq\_7262  
consensus size: 566bp; fragments: 8996; full length: 1 (>=509.4bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

