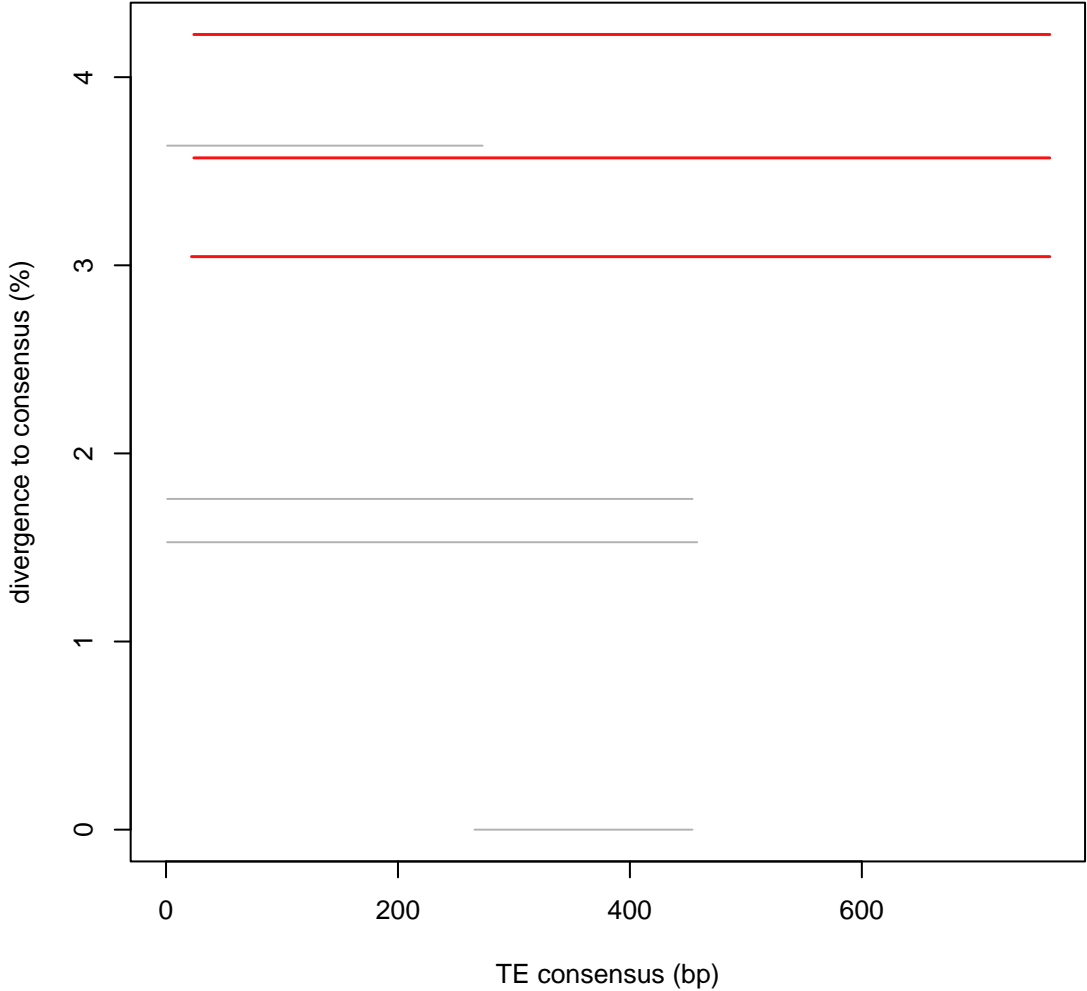
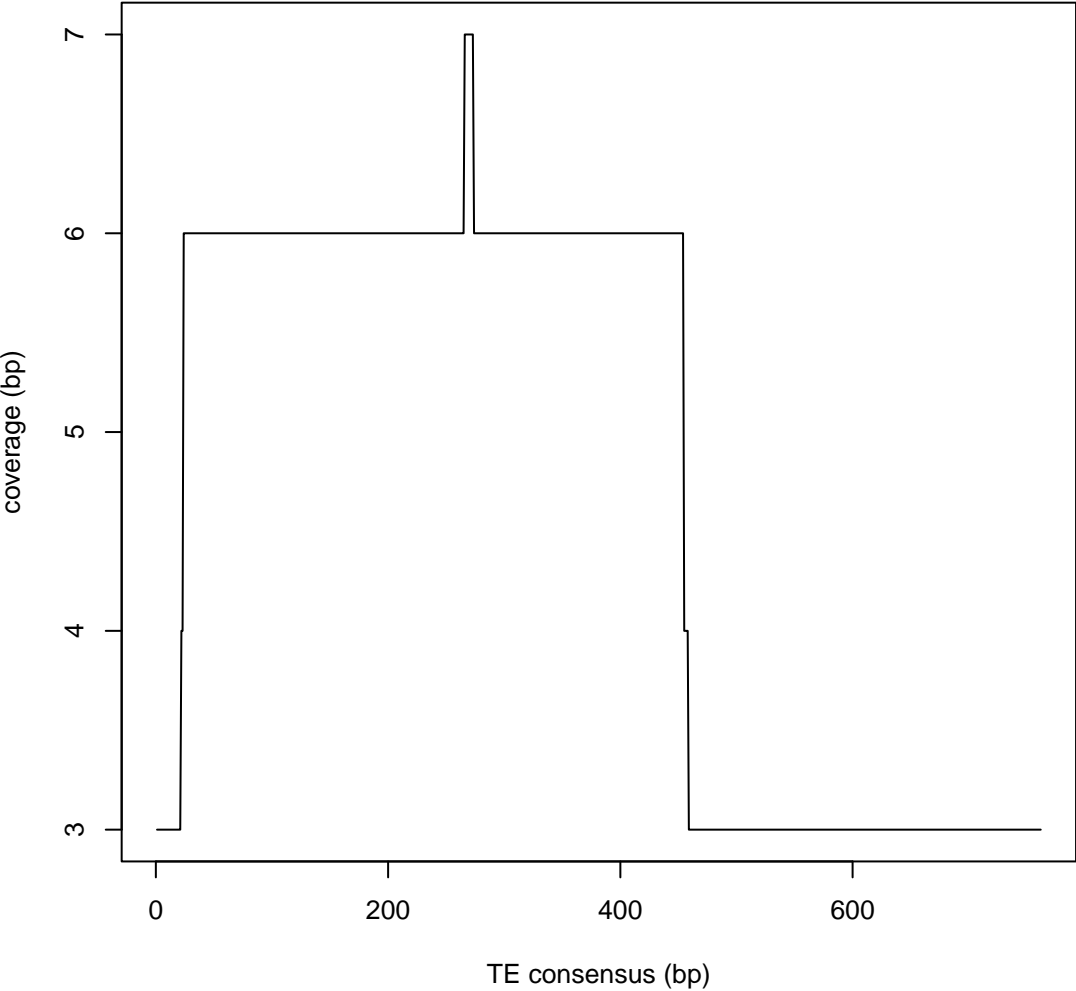


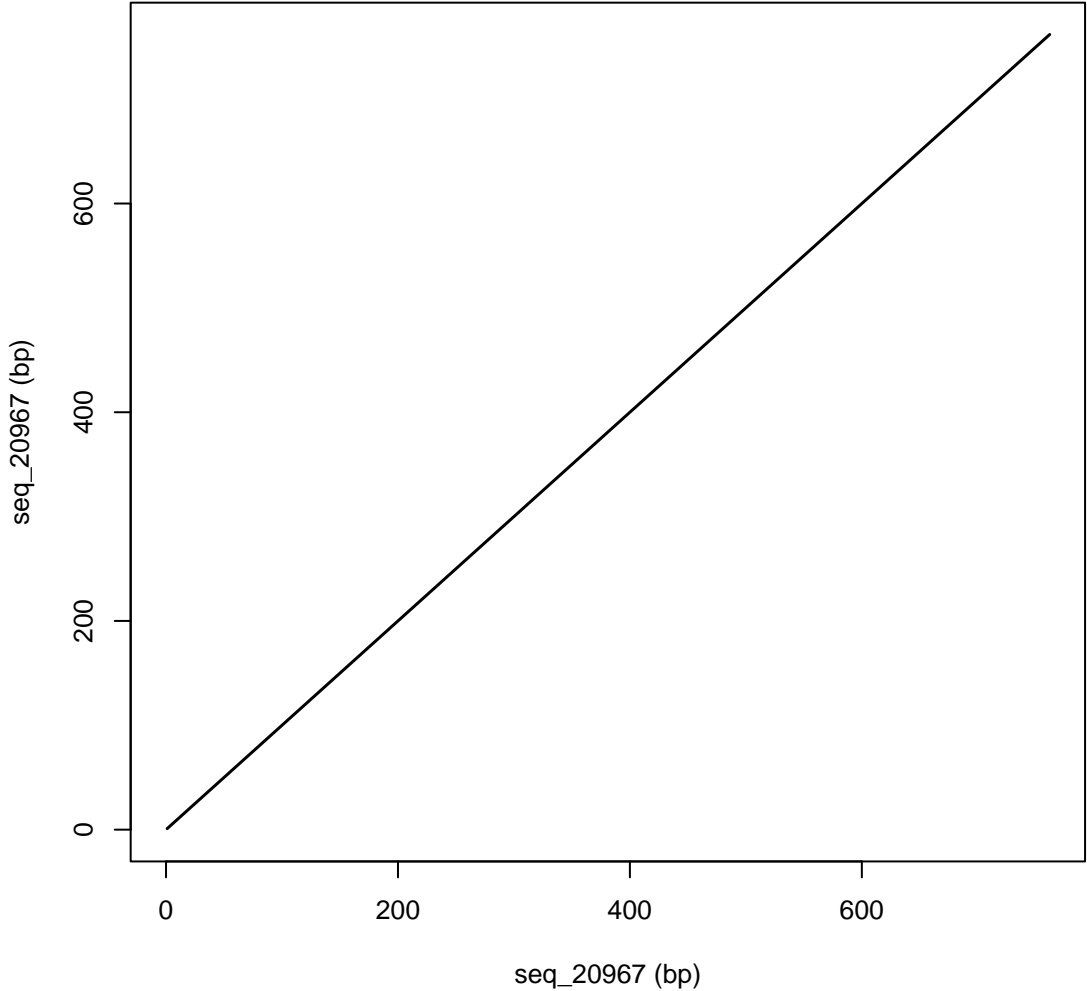
TE: seq\_20967  
consensus size: 762bp; fragments: 7; full length: 3 (>=685.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

