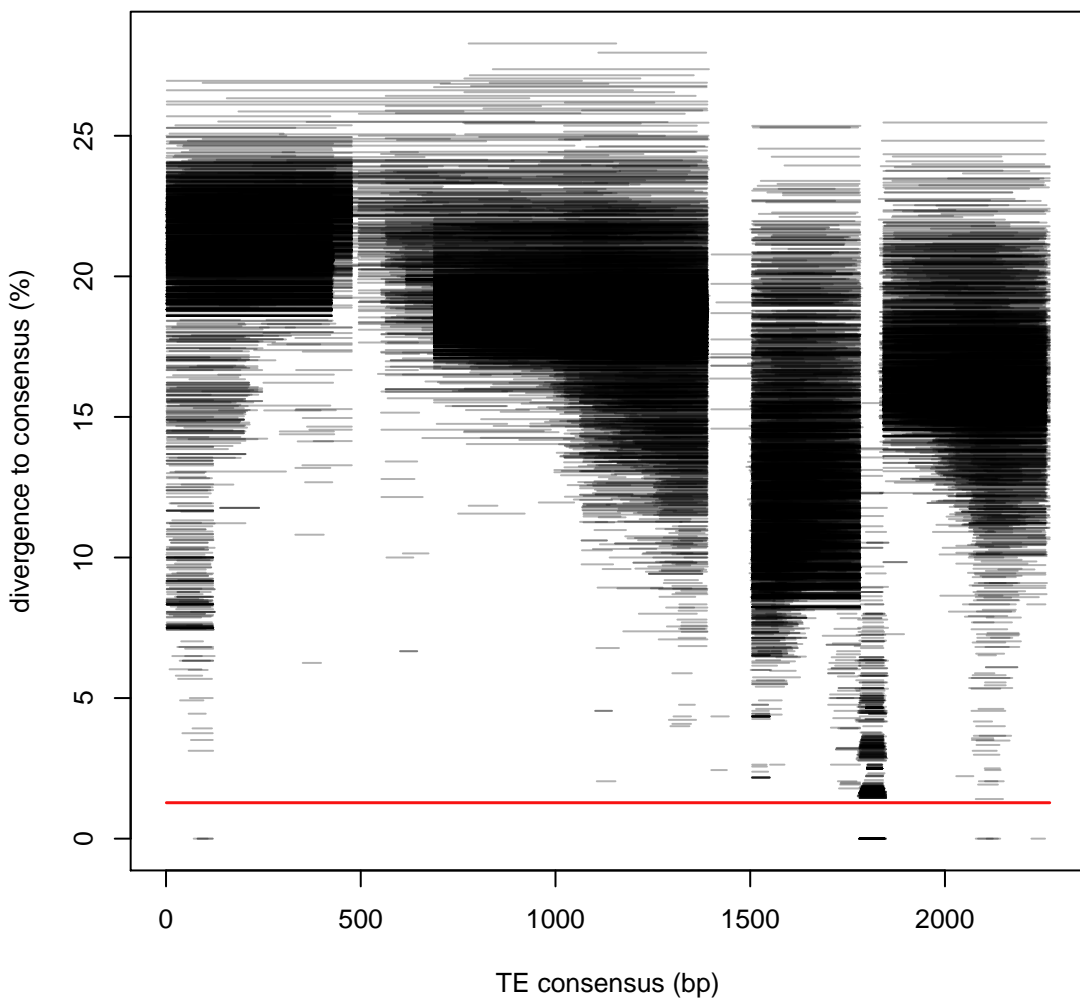
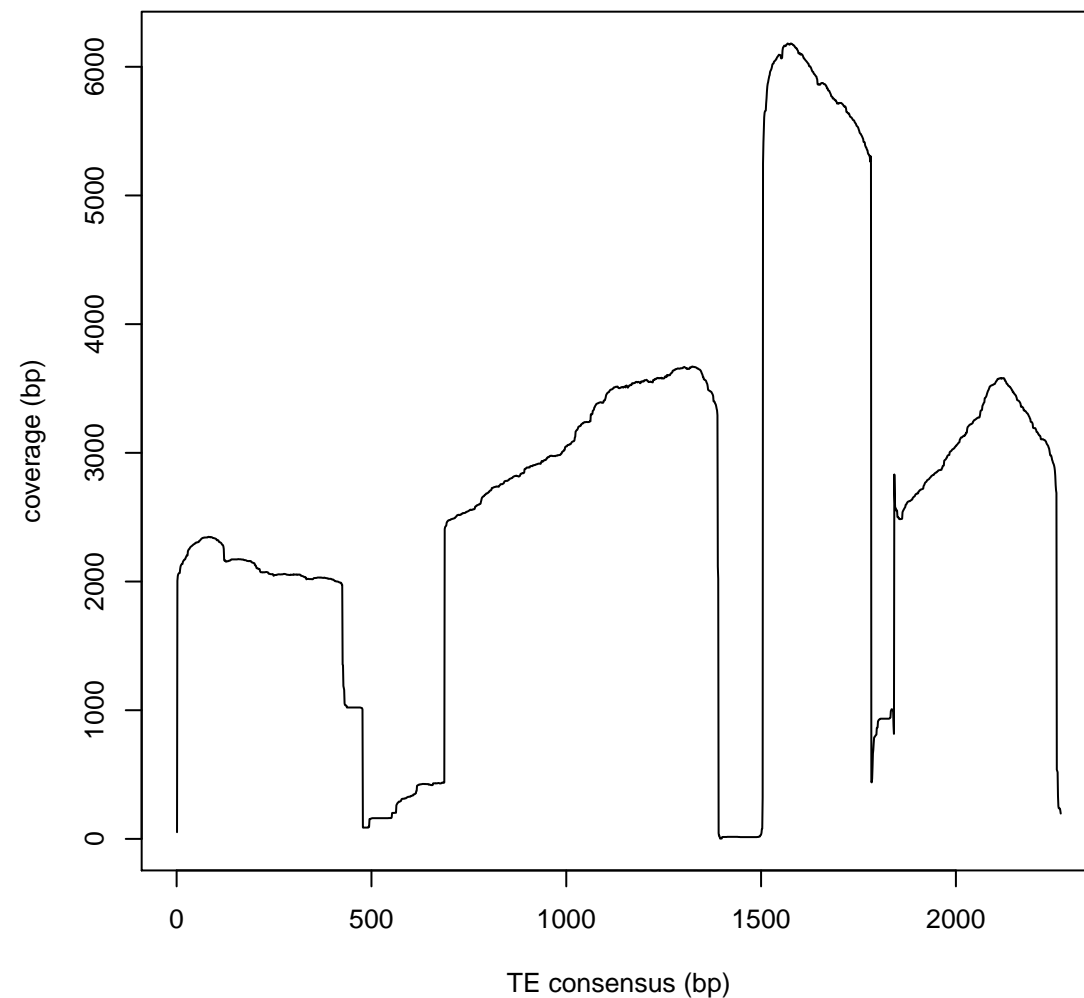


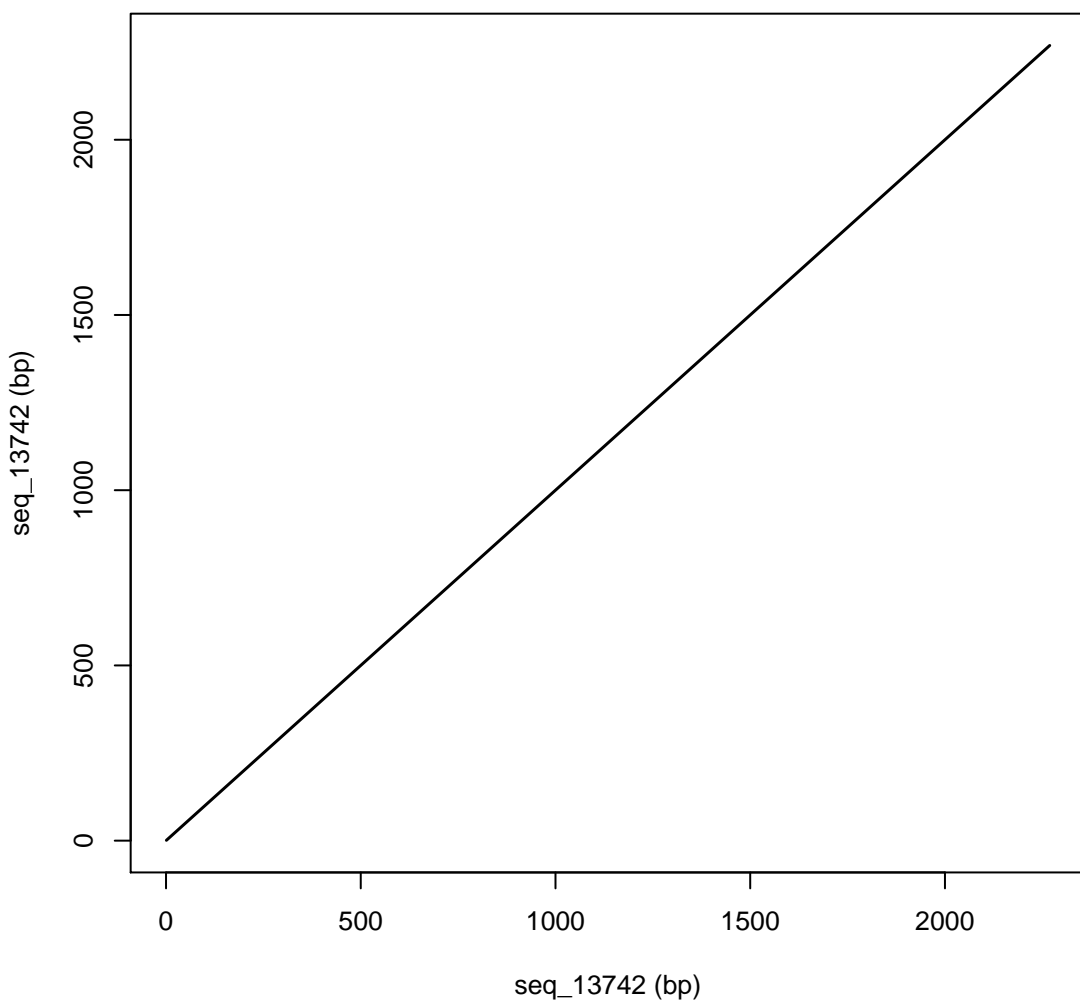
TE: seq\_13742  
consensus size: 2269bp; fragments: 17496; full length: 1 (>=2042.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

