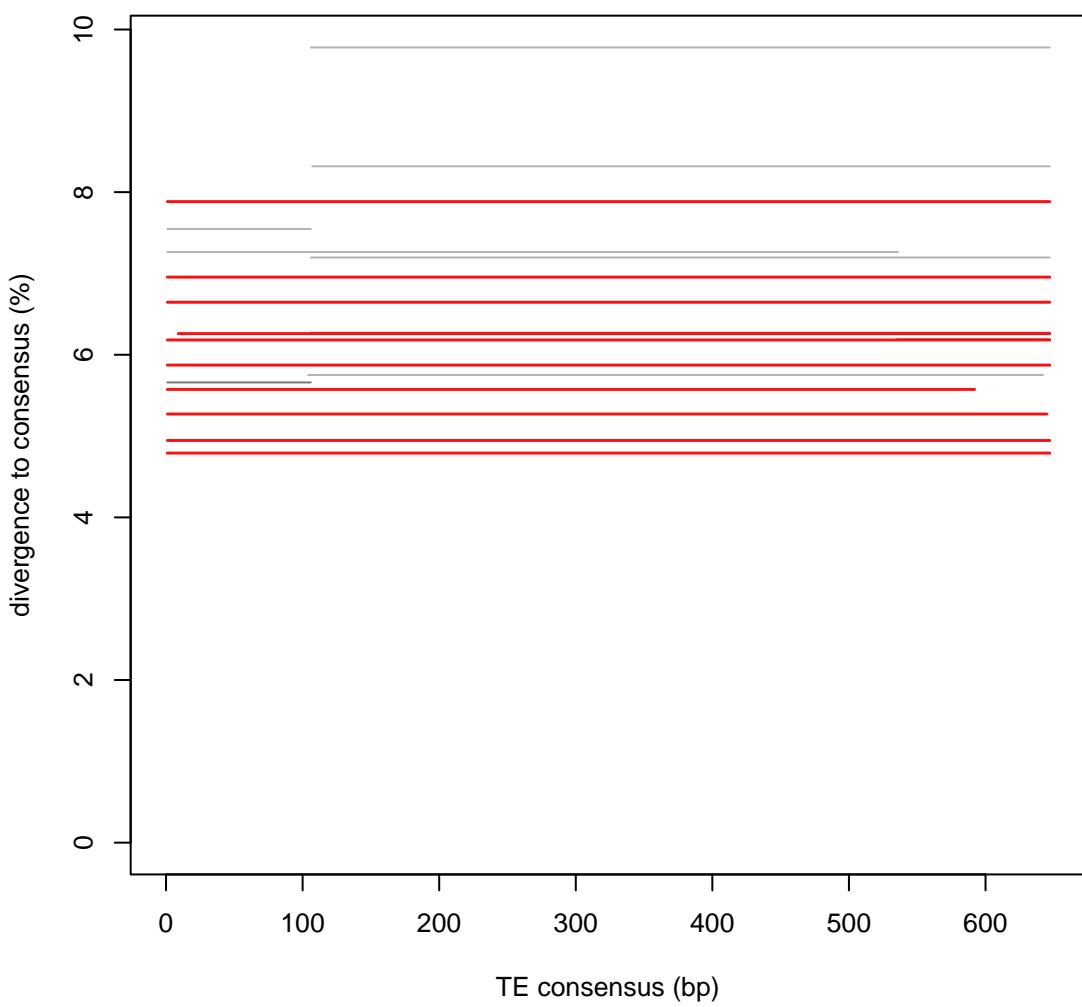
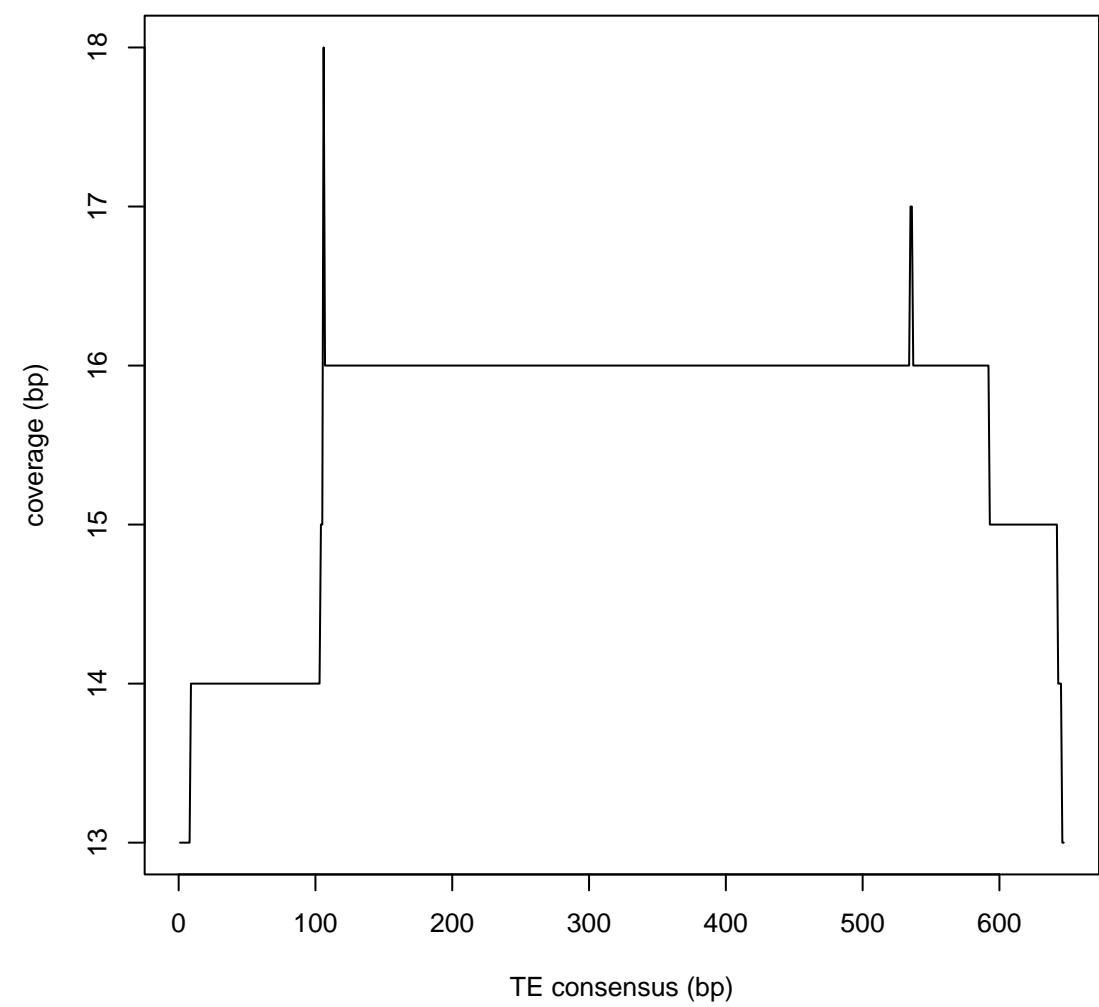


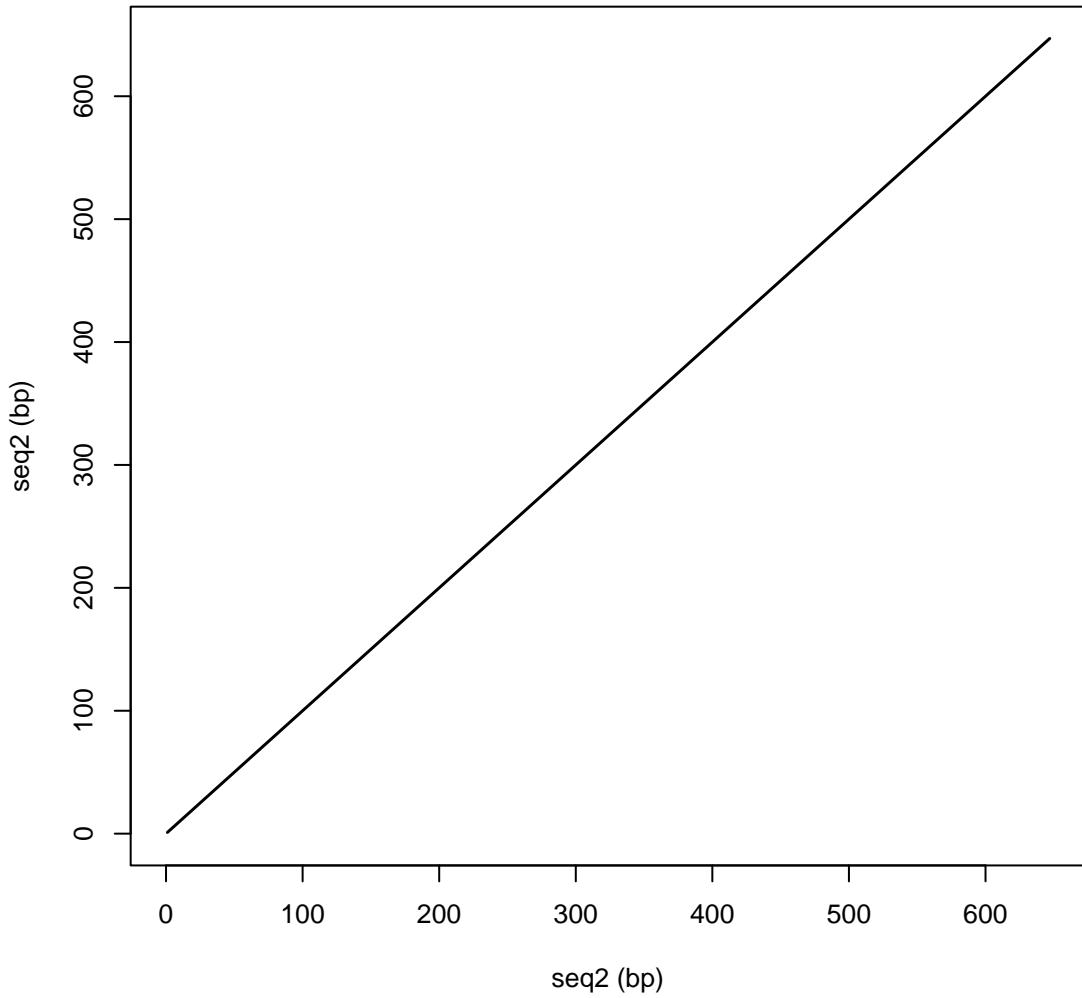
TE: seq2
consensus size: 647bp; fragments: 20; full length: 10 (≥ 582.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

