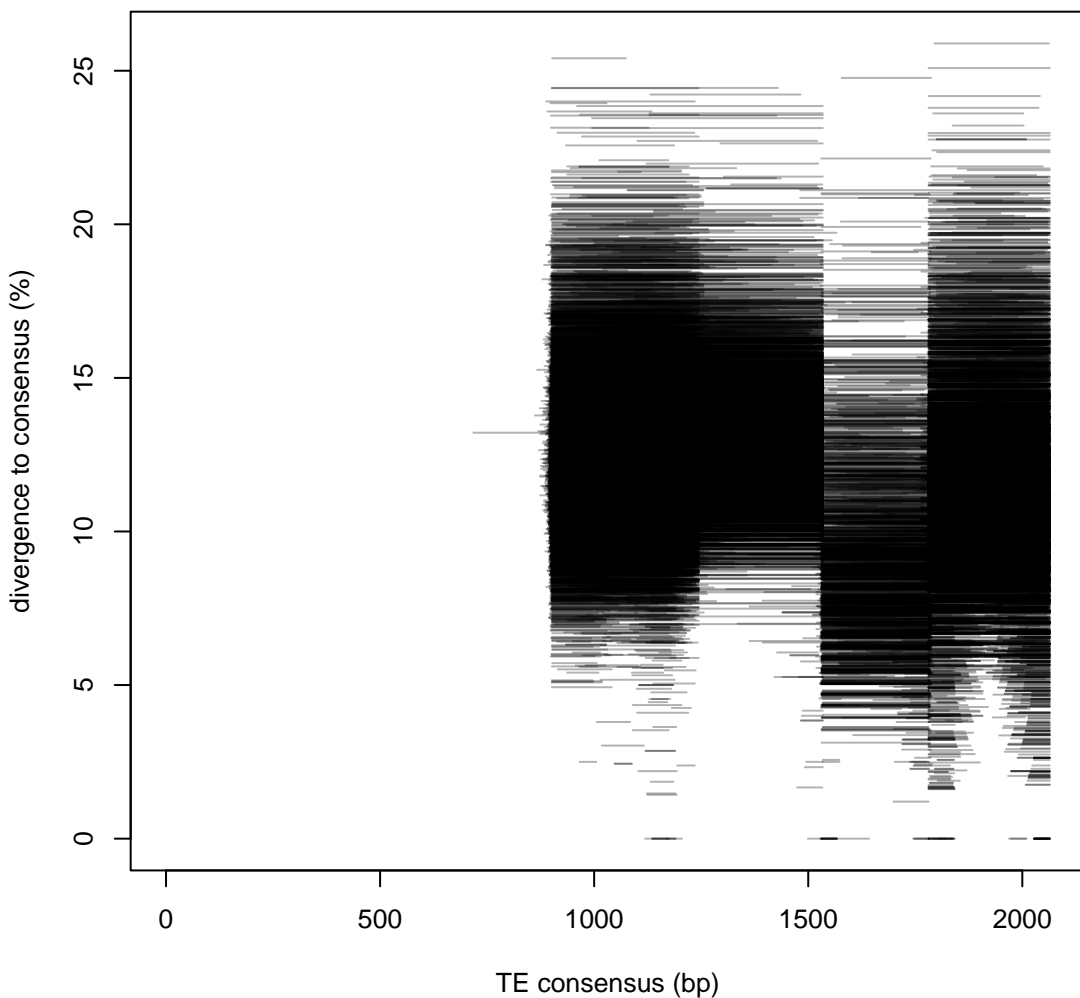
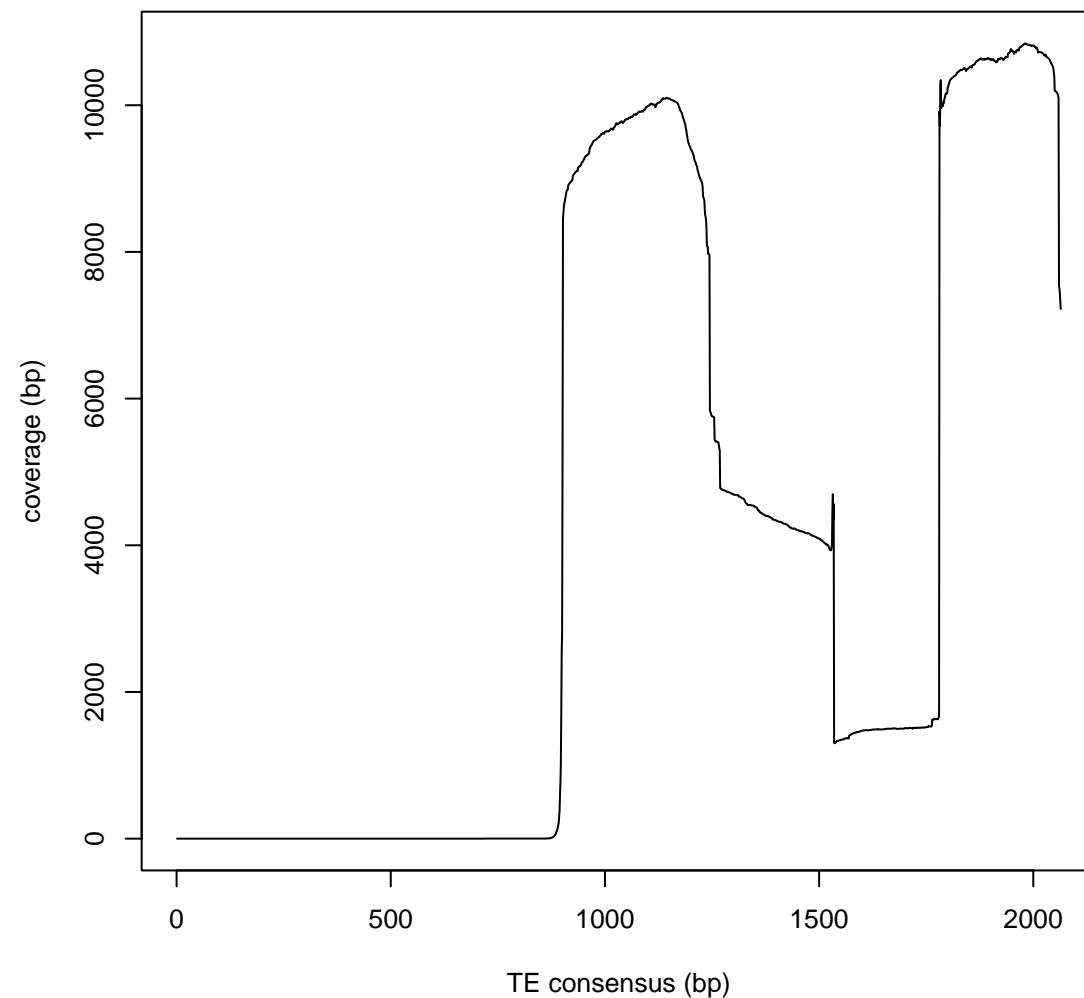


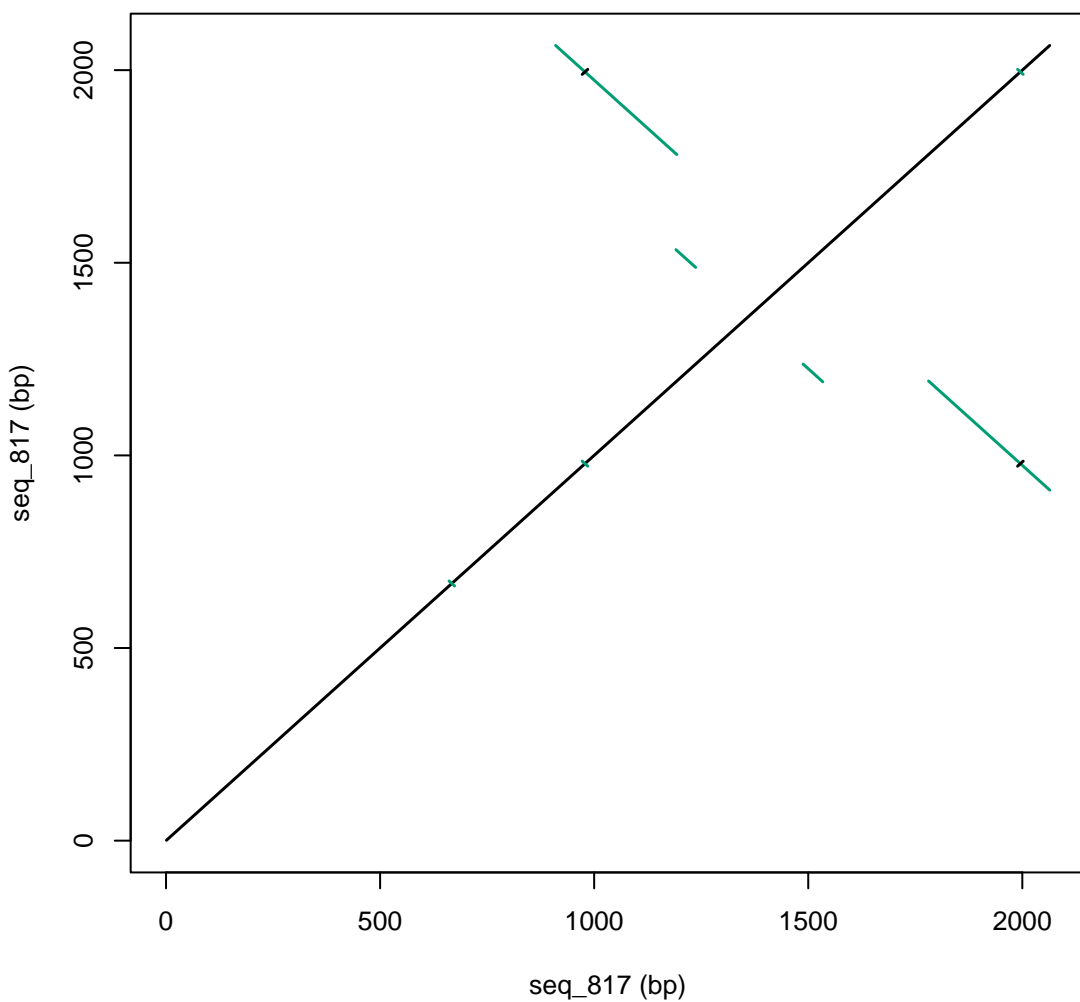
TE: seq\_817  
consensus size: 2064bp; fragments: 24305; full length: 0 (>=1857.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

