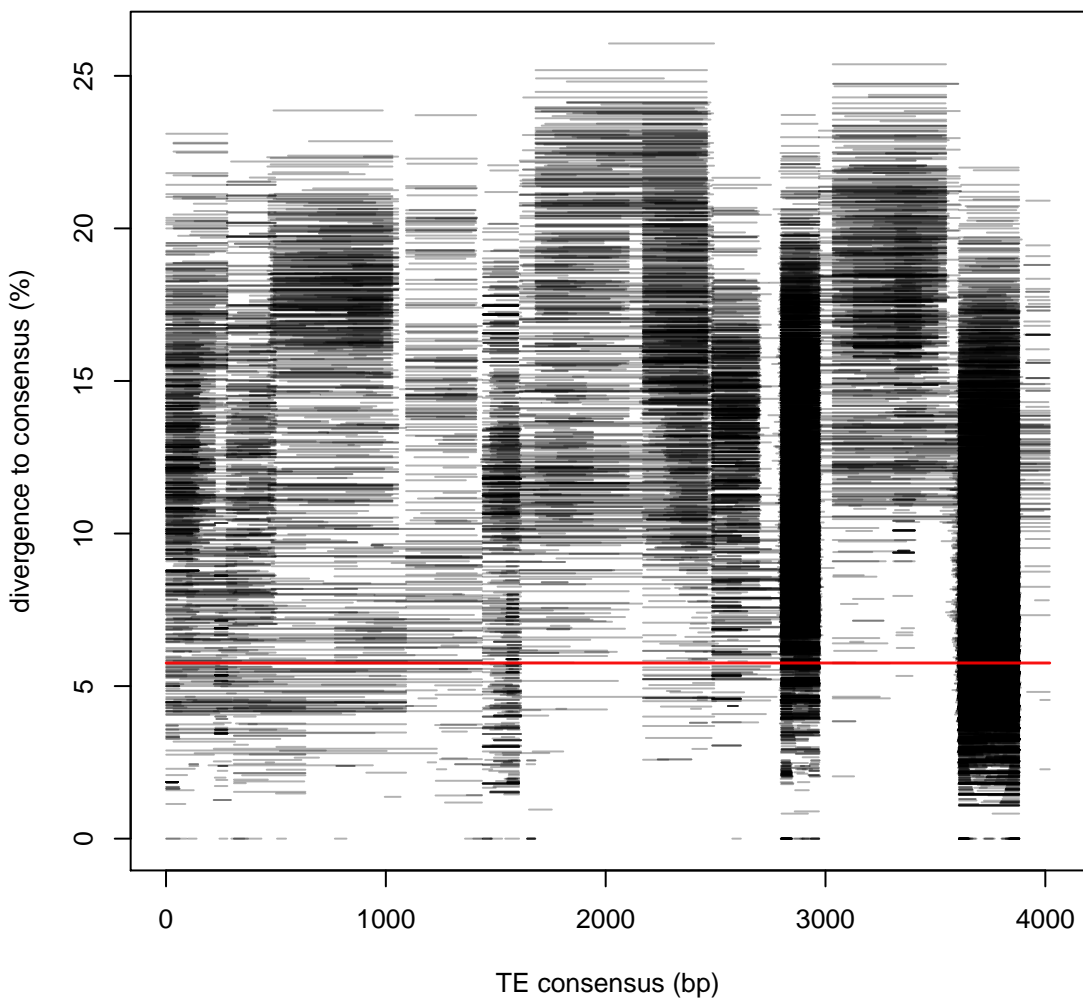
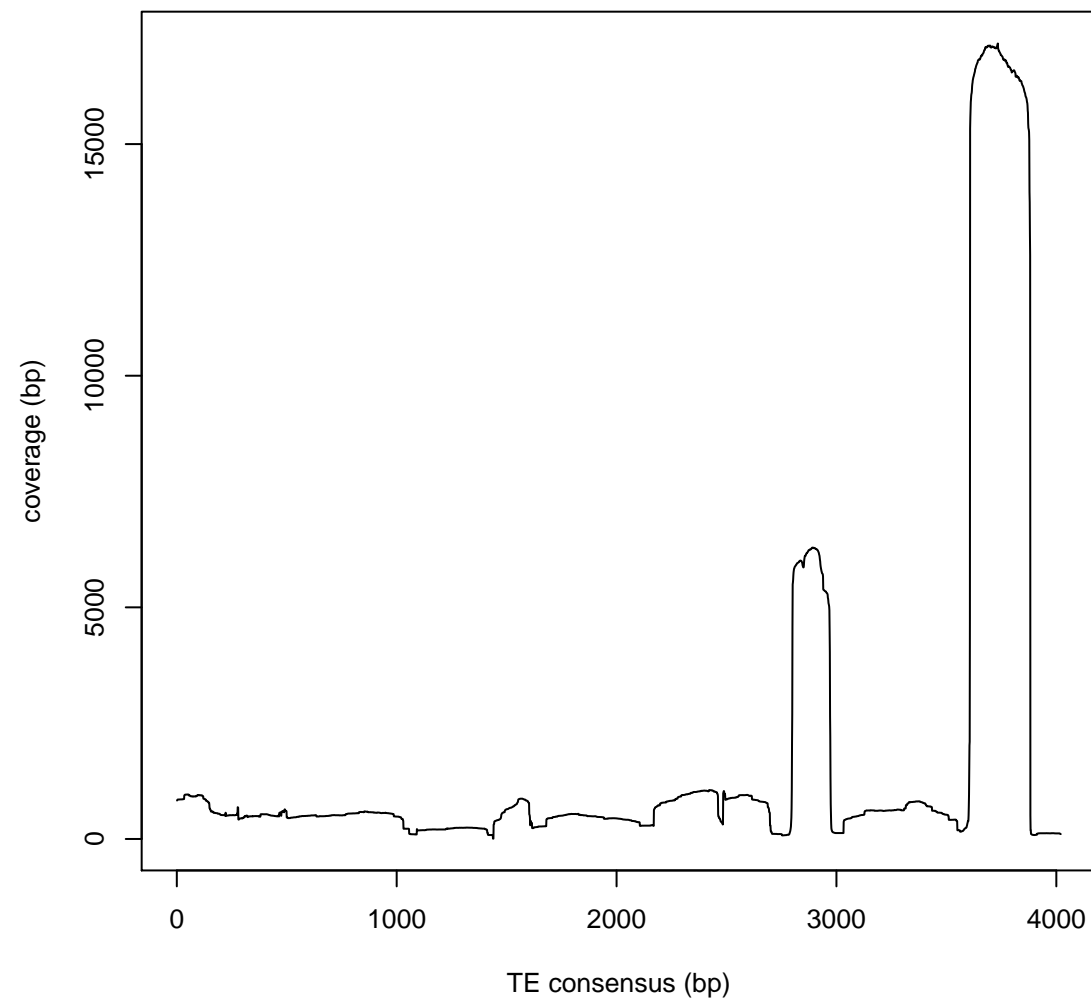


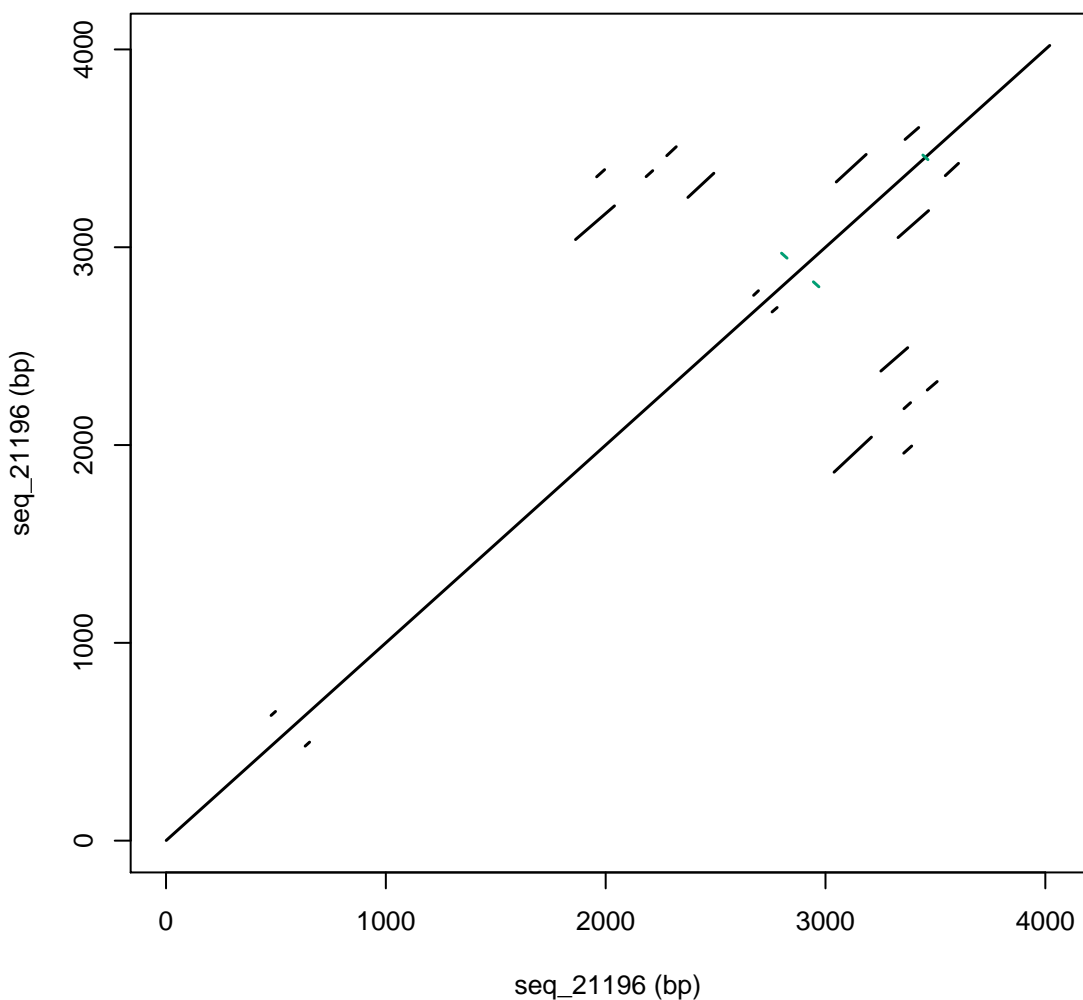
TE: seq\_21196  
consensus size: 4020bp; fragments: 32054; full length: 1 (>=3618bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

