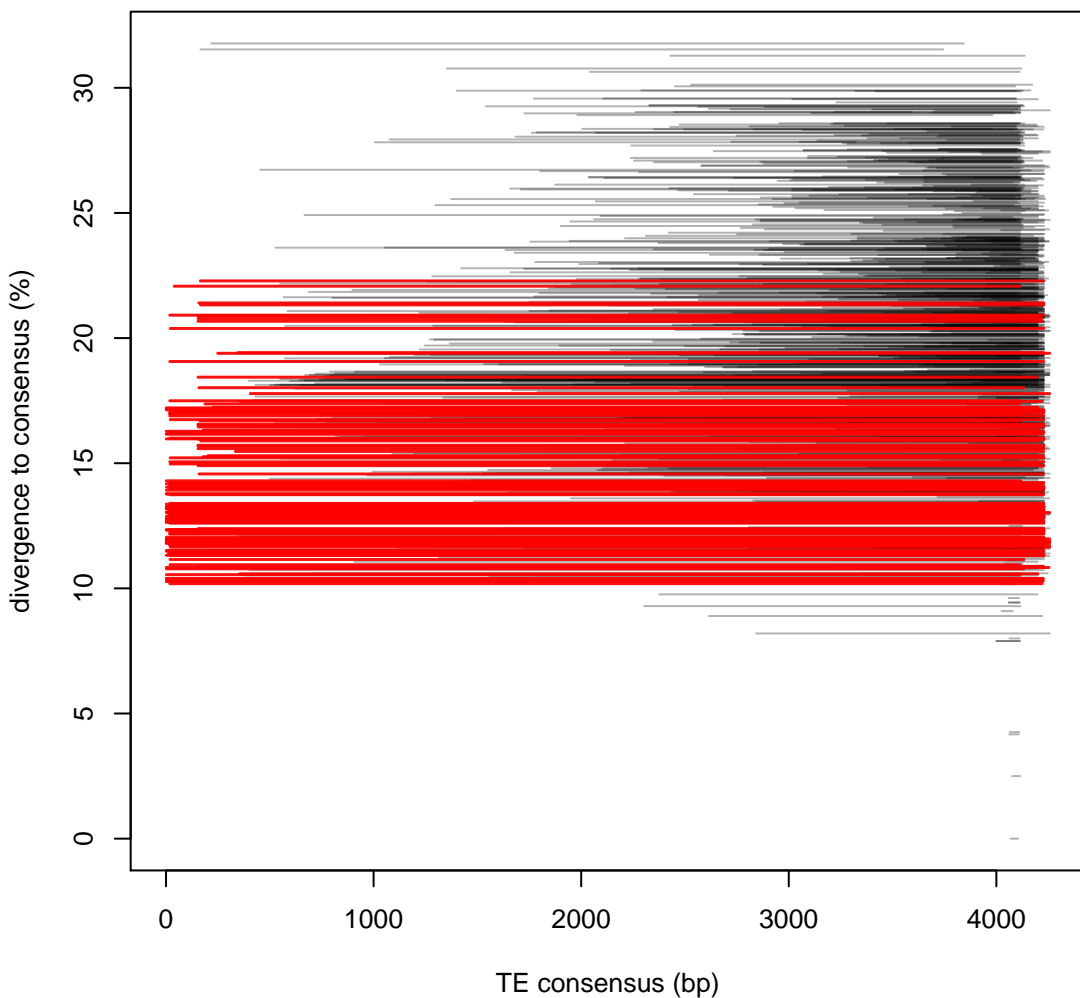
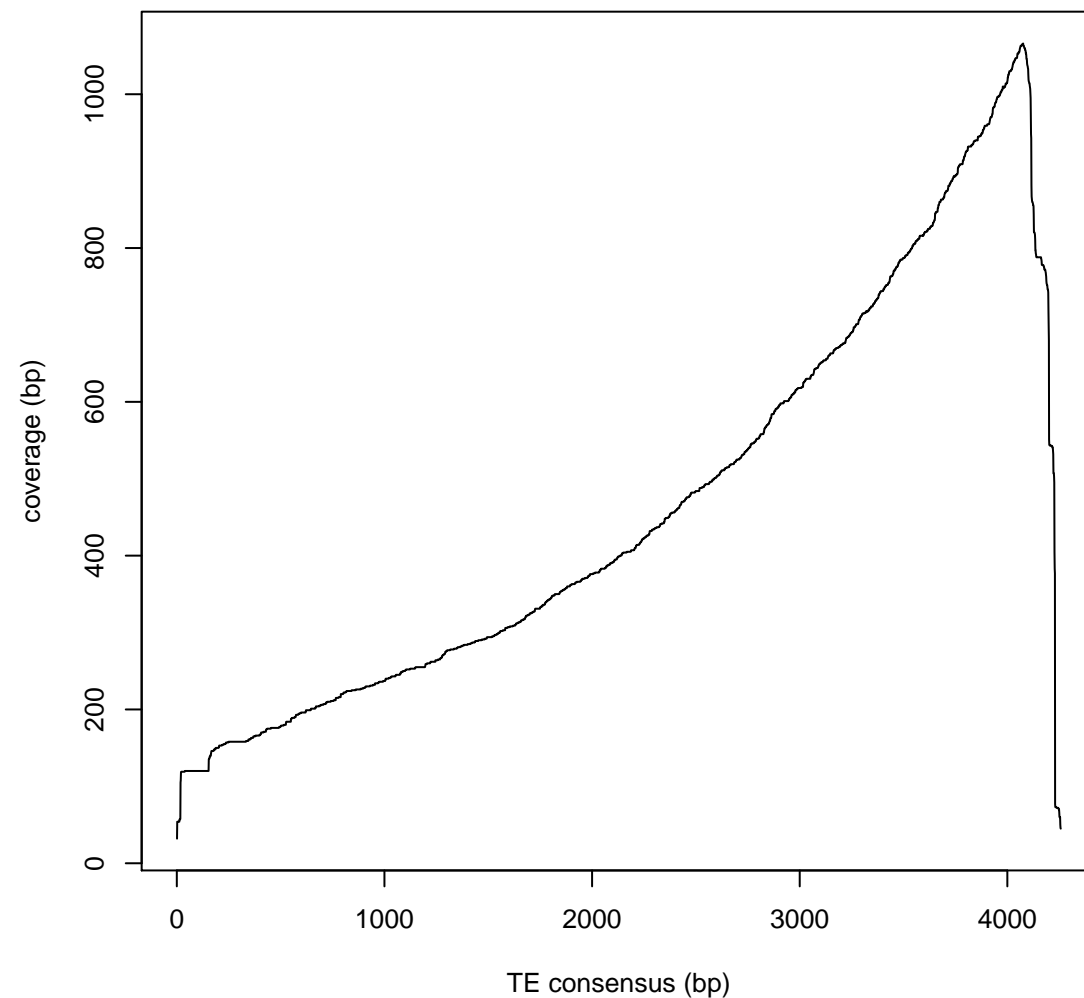


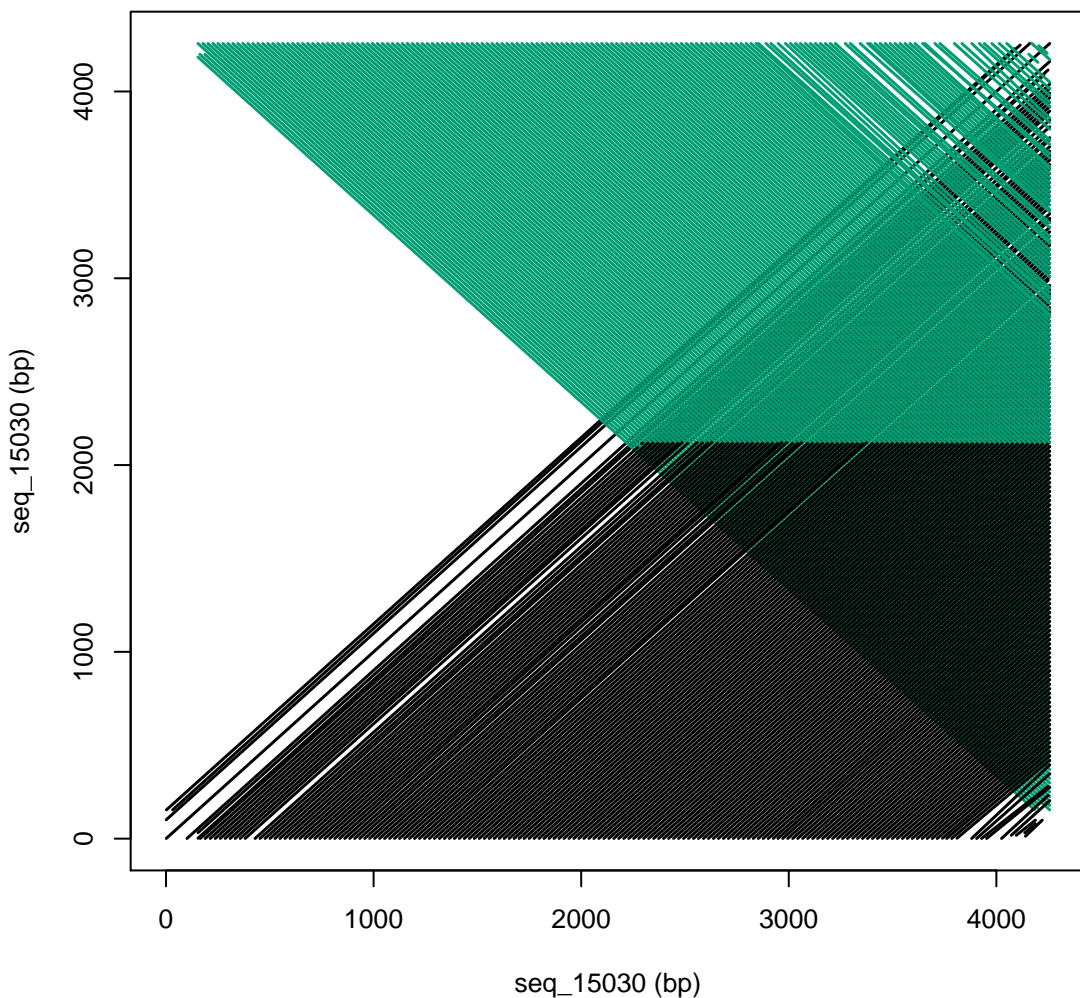
TE: seq\_15030  
consensus size: 4257bp; fragments: 1073; full length: 162 (>=3831.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

