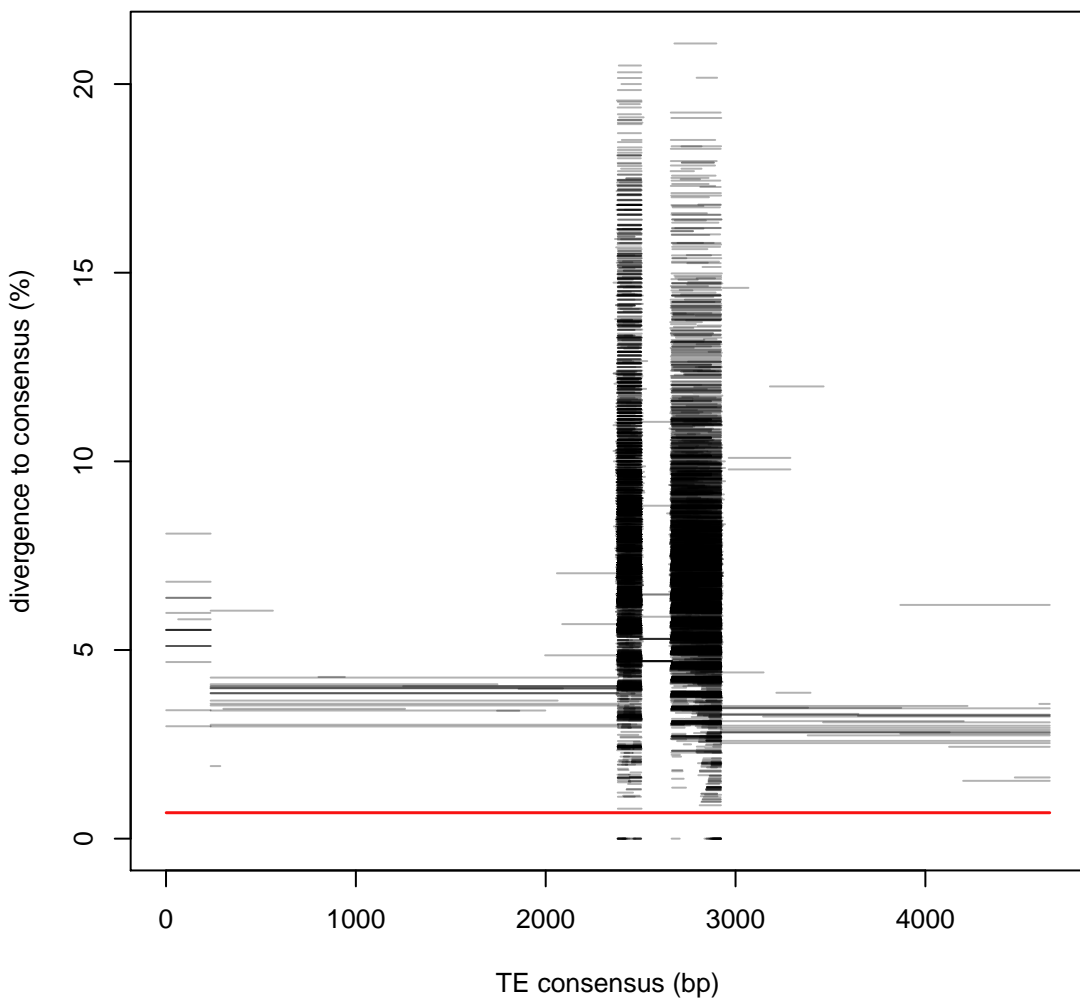
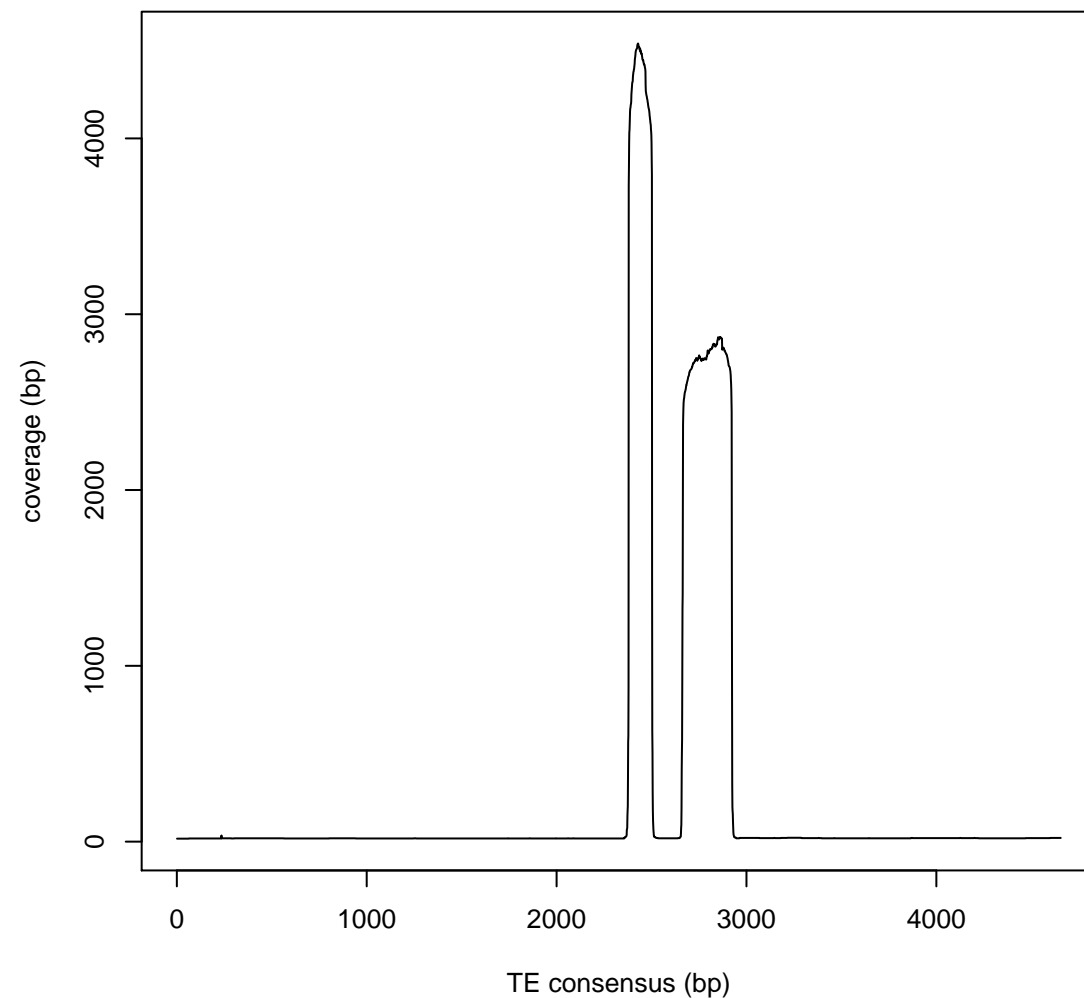


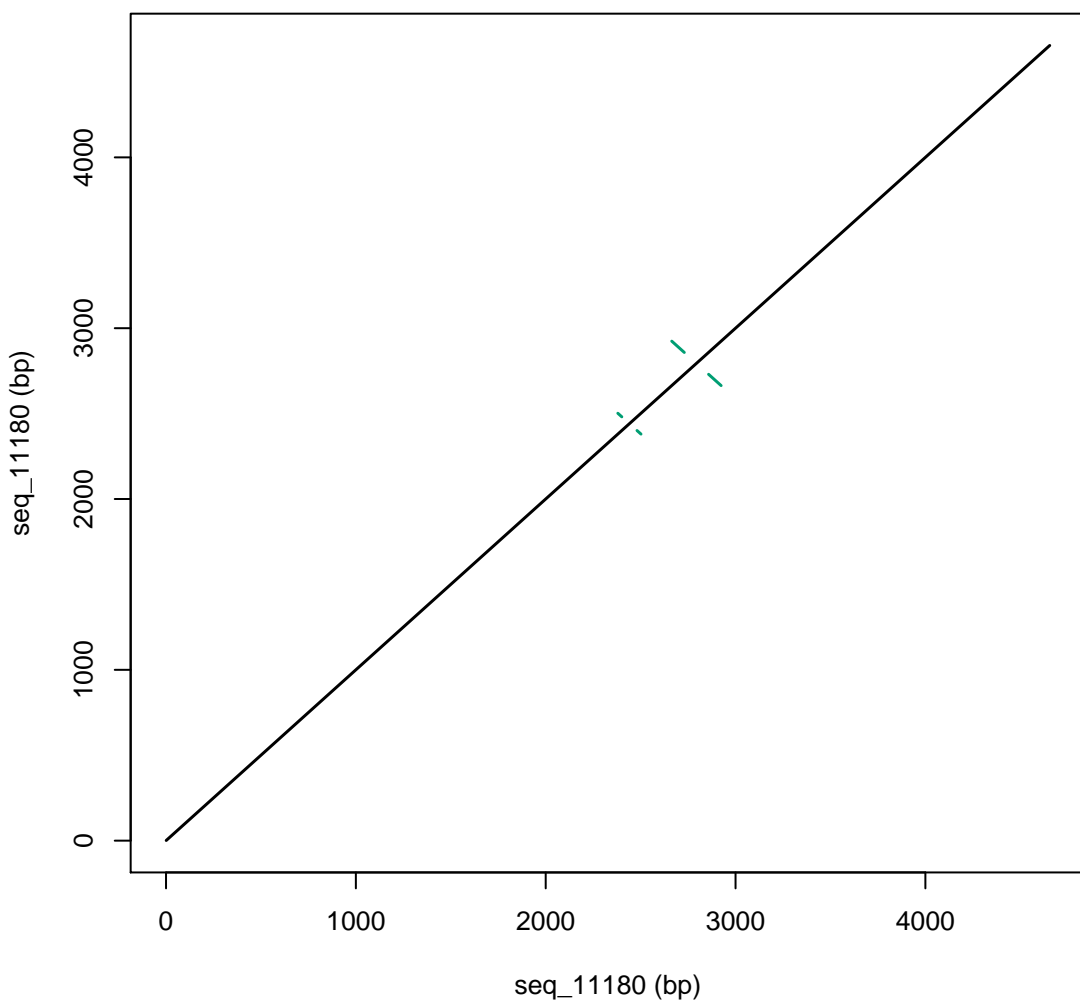
TE: seq\_11180  
consensus size: 4655bp; fragments: 7988; full length: 1 (>=4189.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

