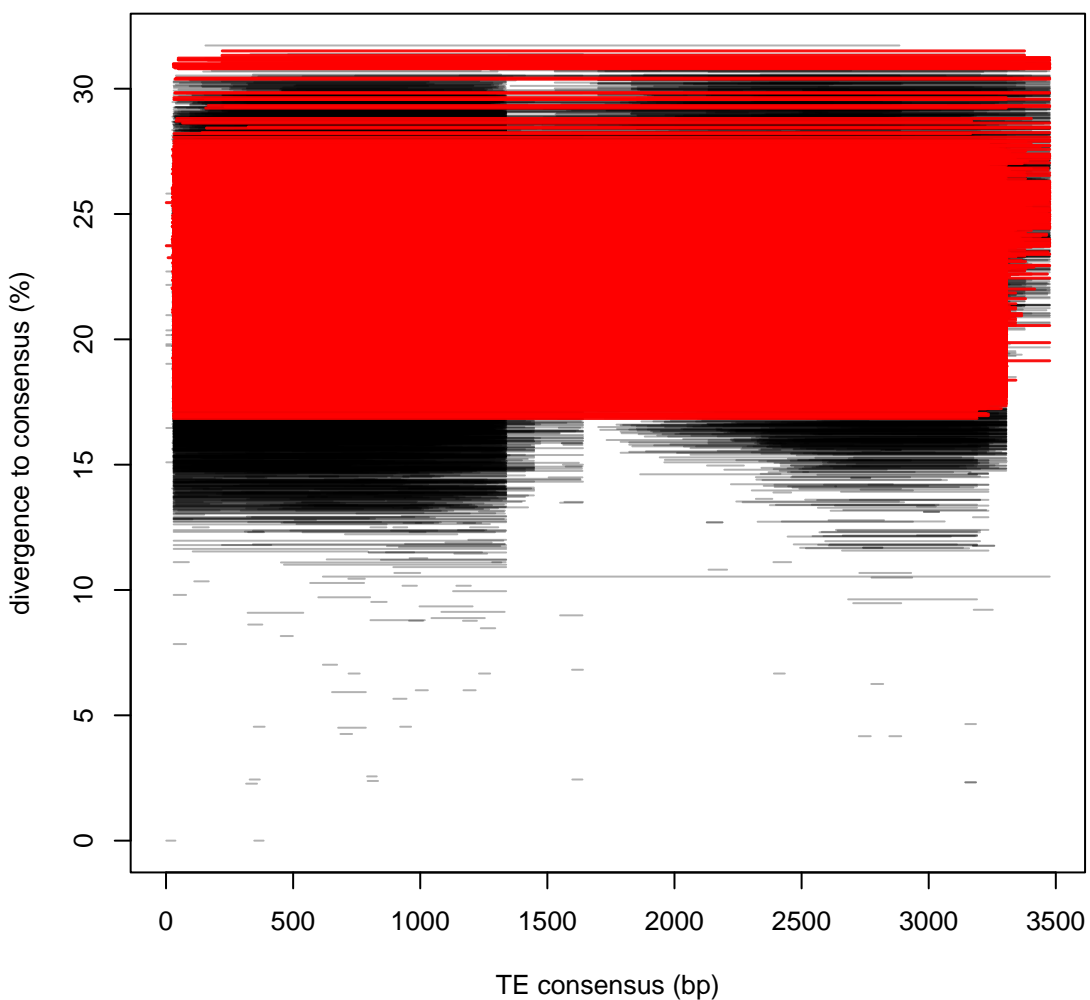
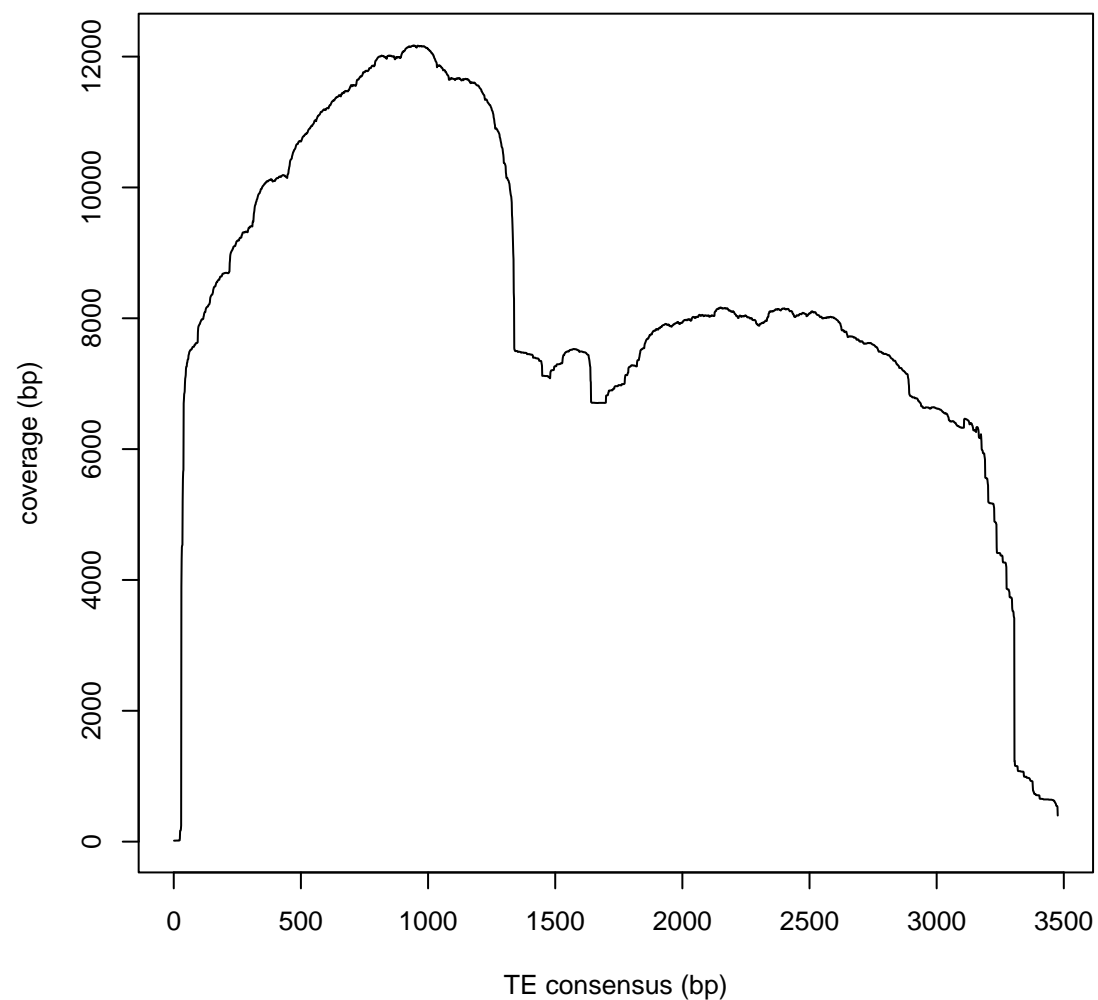


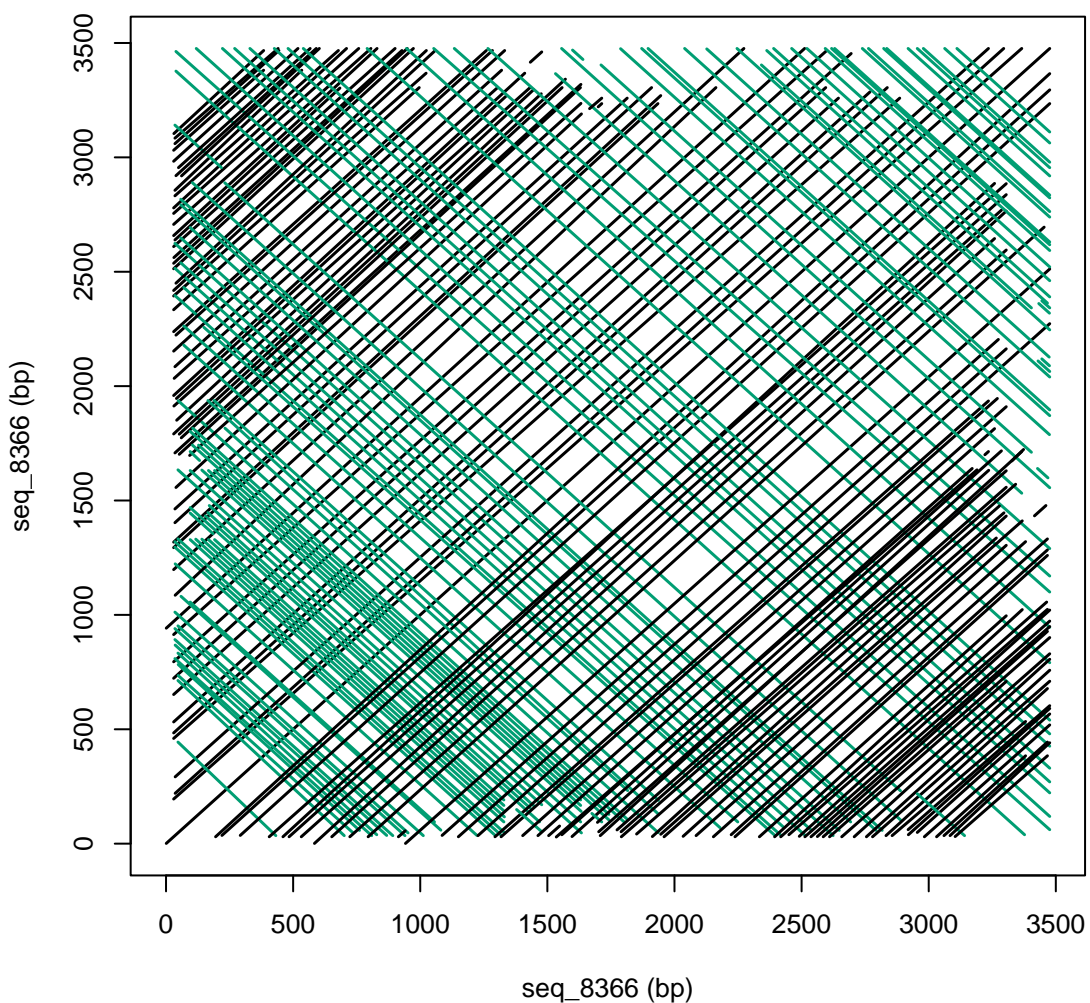
TE: seq\_8366  
consensus size: 3476bp; fragments: 18102; full length: 2456 ( $\geq 3128.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

