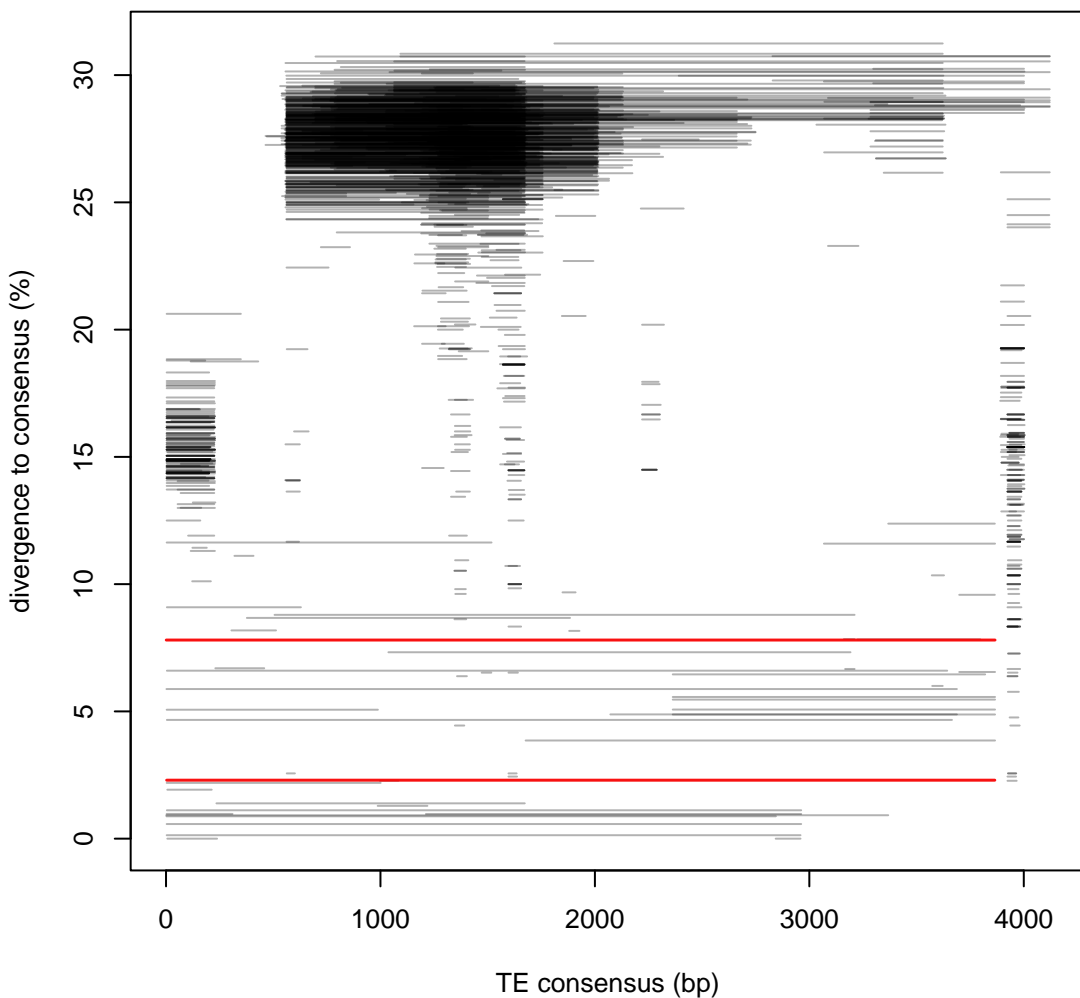
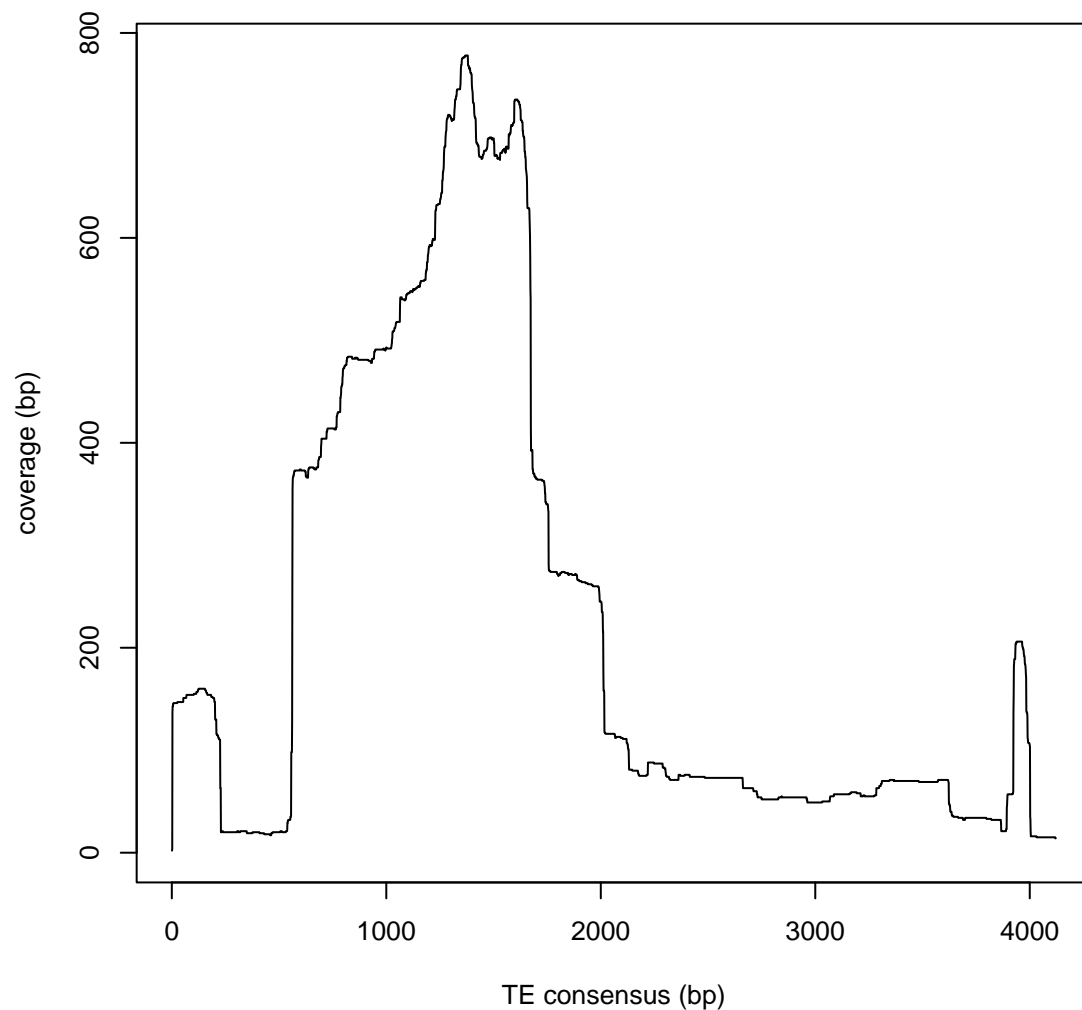


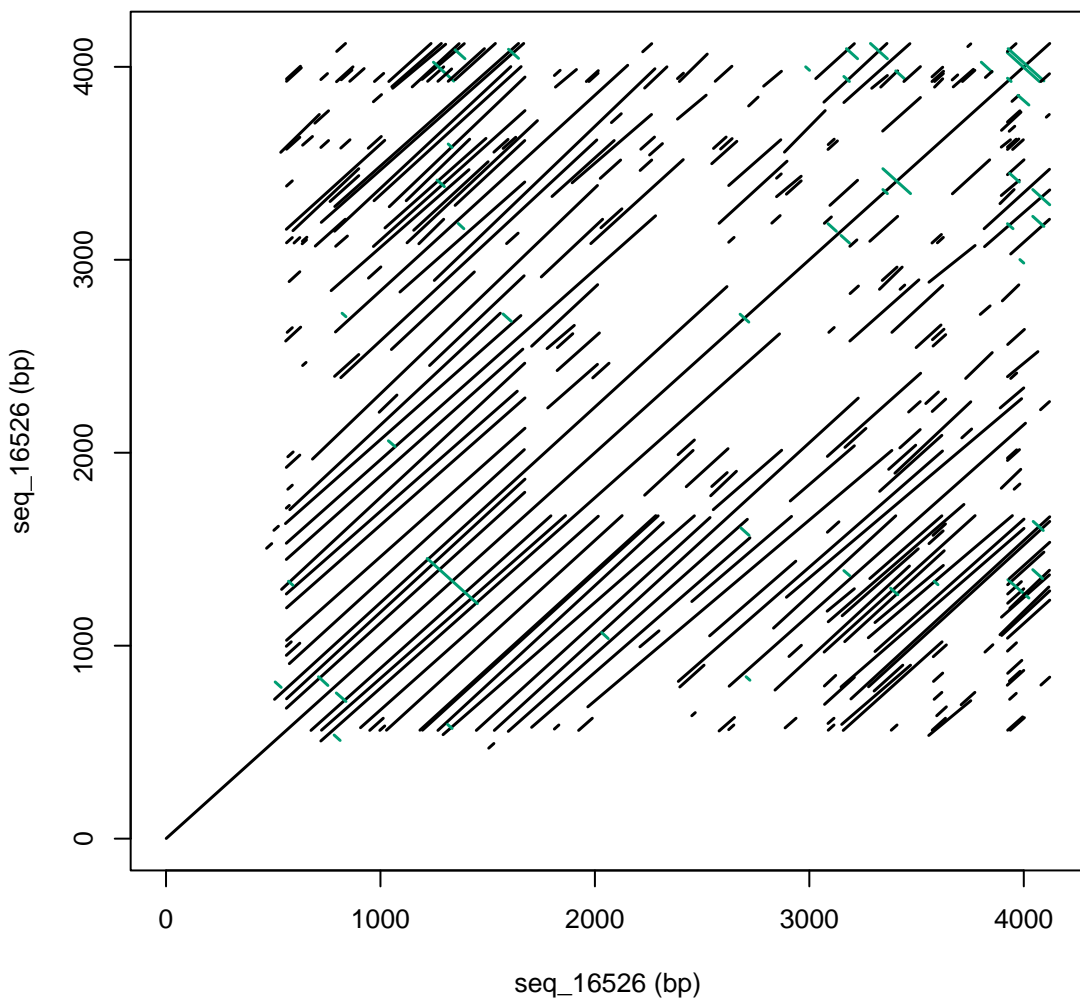
TE: seq\_16526  
consensus size: 4121bp; fragments: 1313; full length: 2 ( $\geq 3708.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

