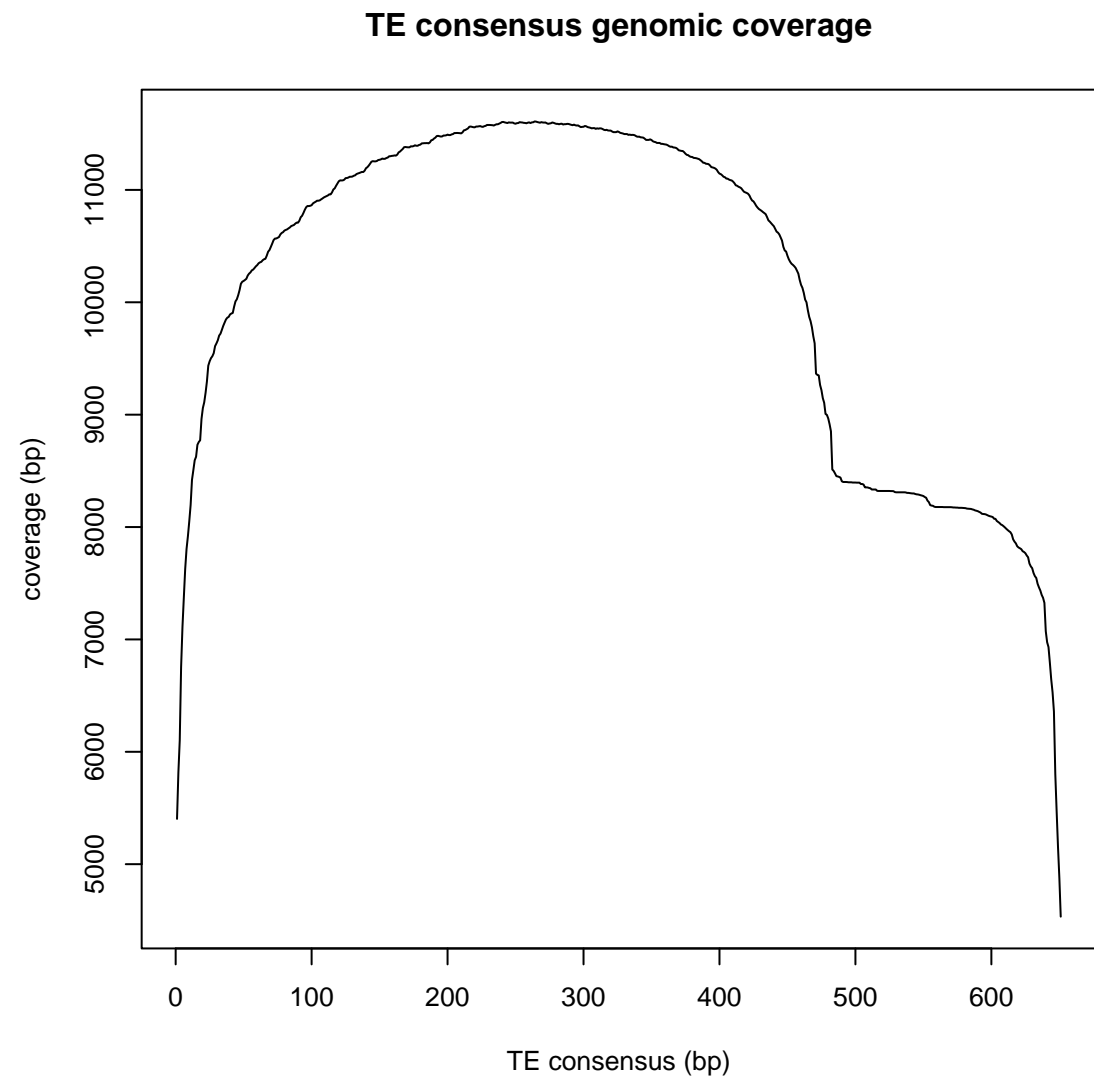
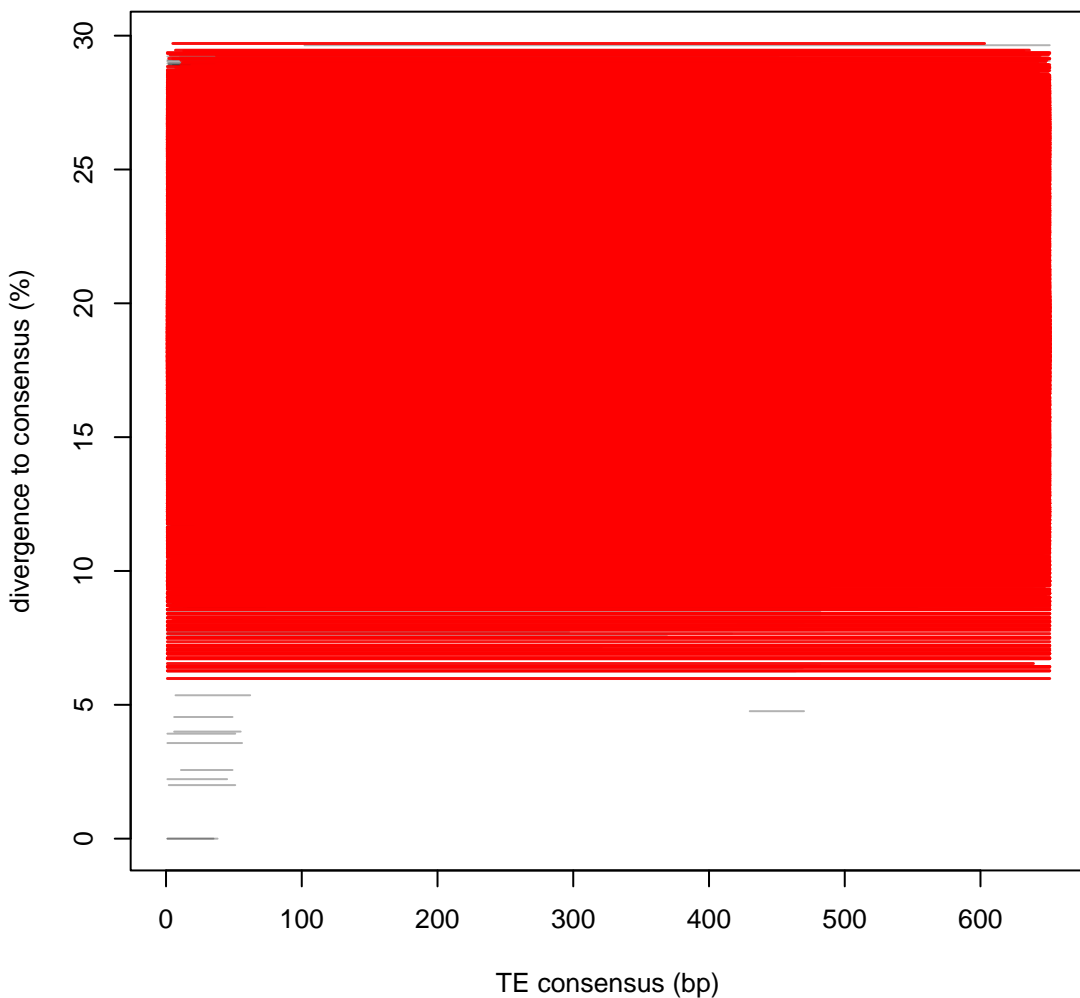
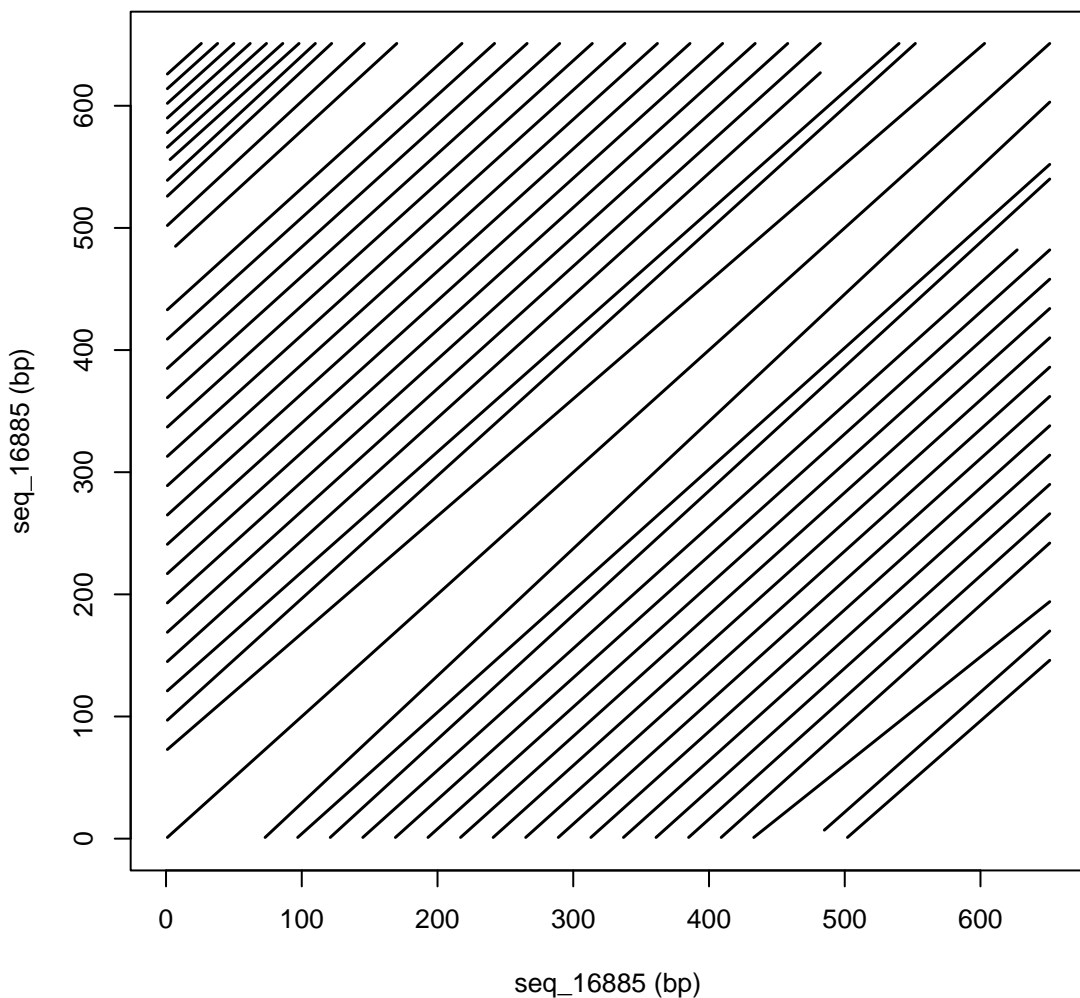


TE: seq\_16885  
consensus size: 651bp; fragments: 12382; full length: 7027 ( $\geq 585.9$ bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

