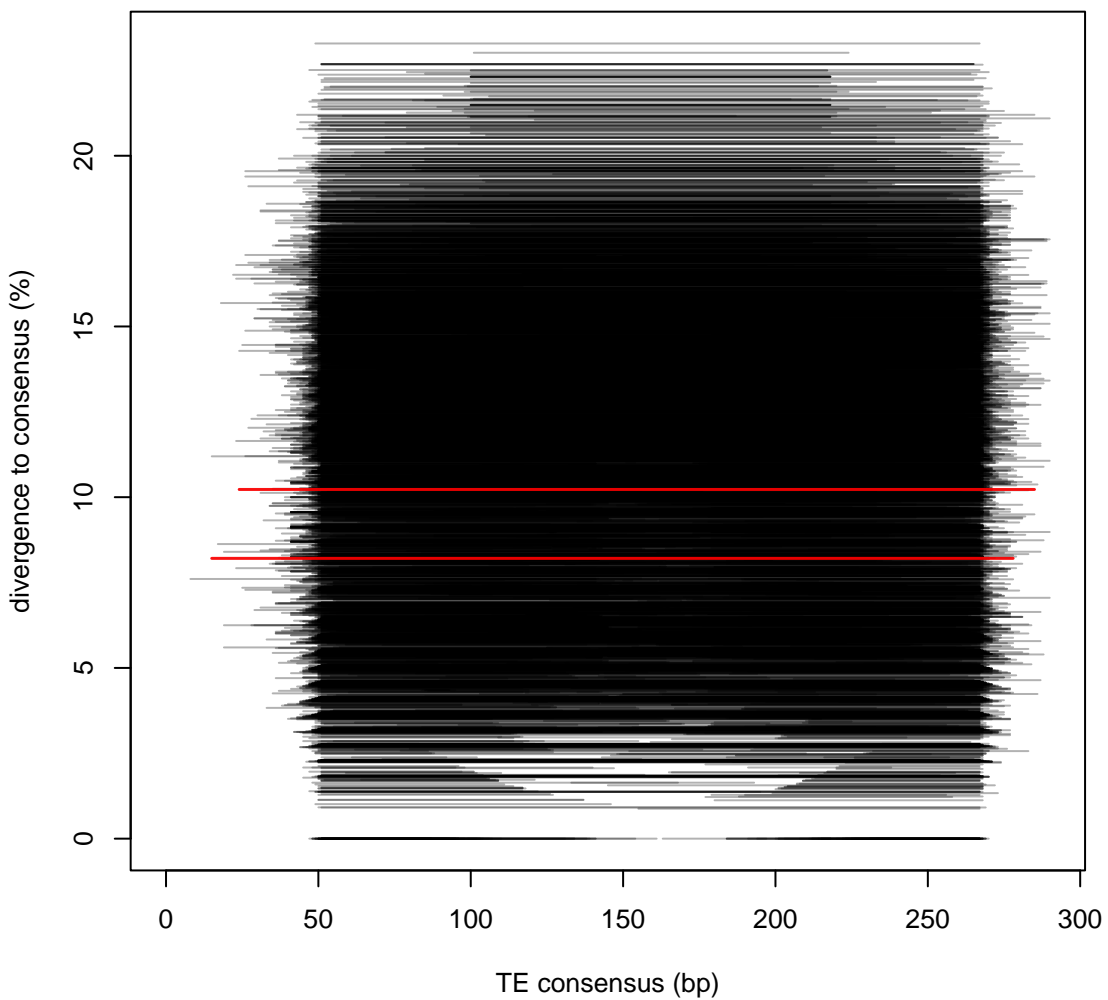
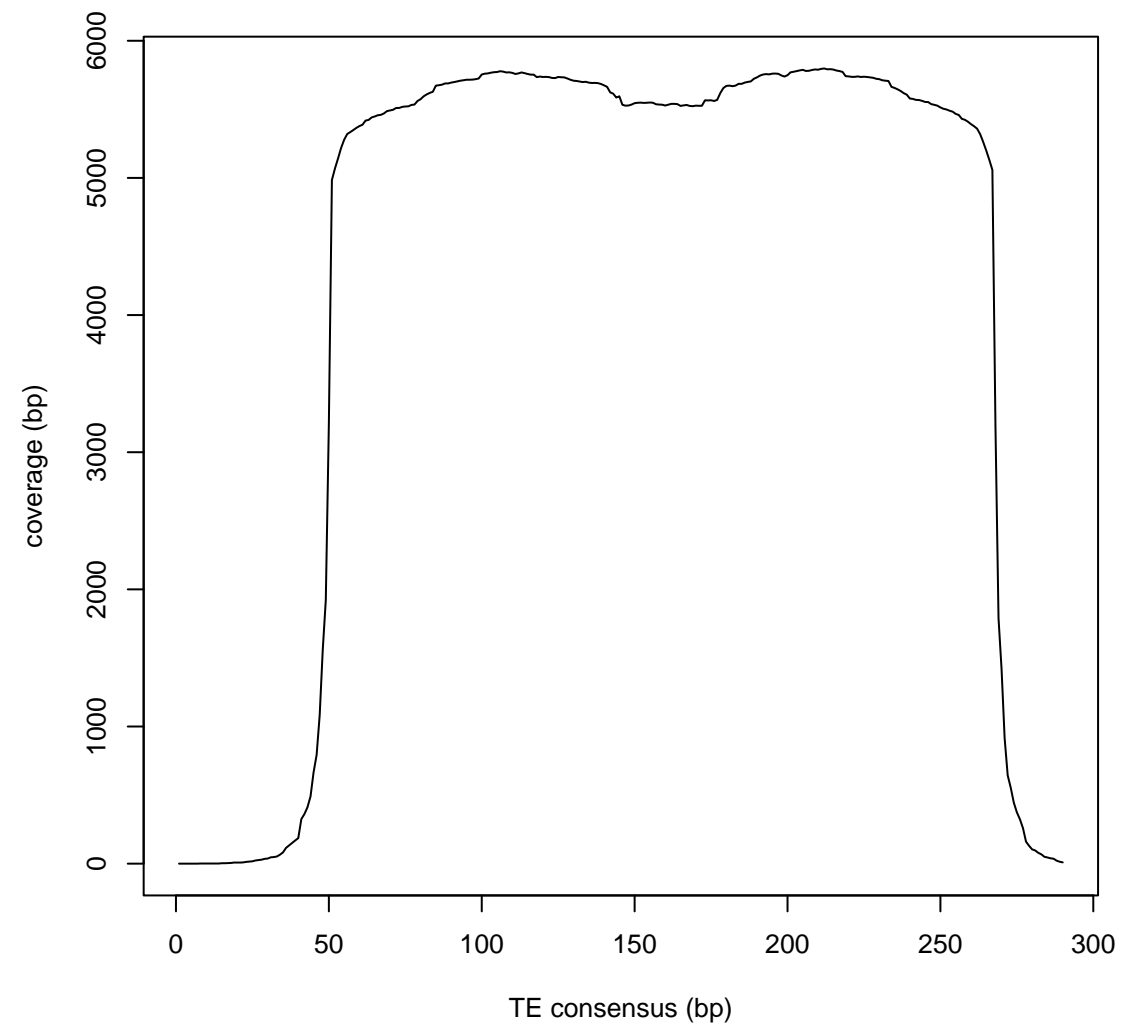


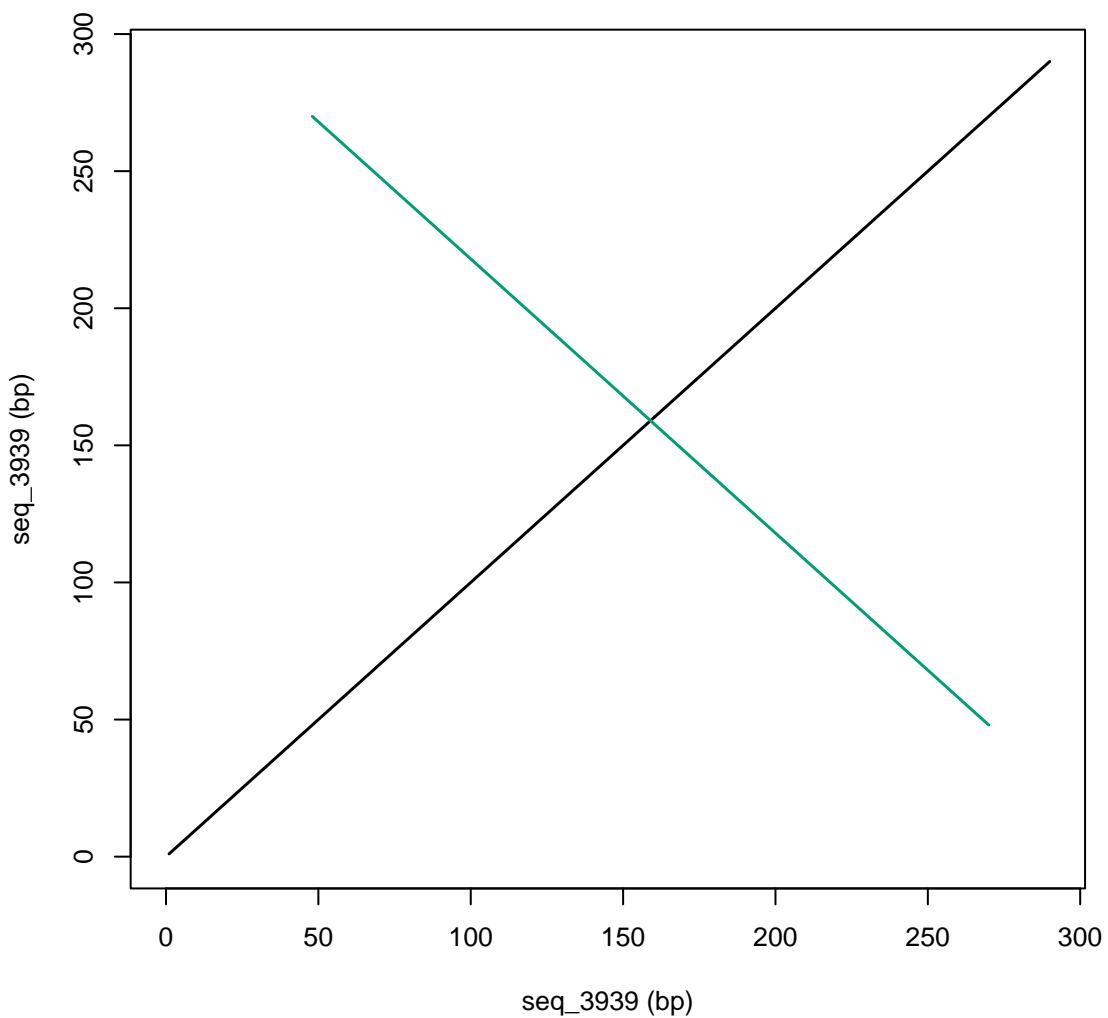
TE: seq\_3939  
consensus size: 290bp; fragments: 6574; full length: 2 ( $\geq 261$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

