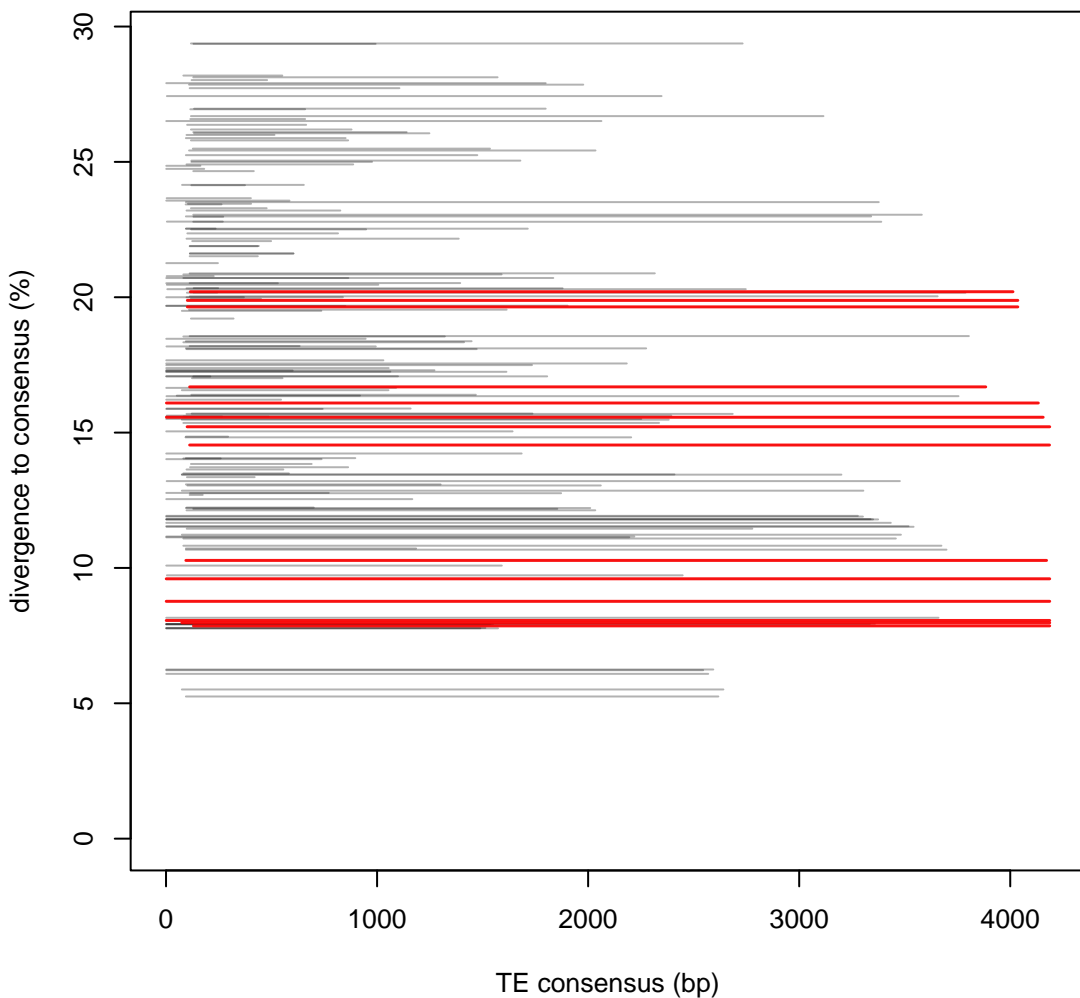
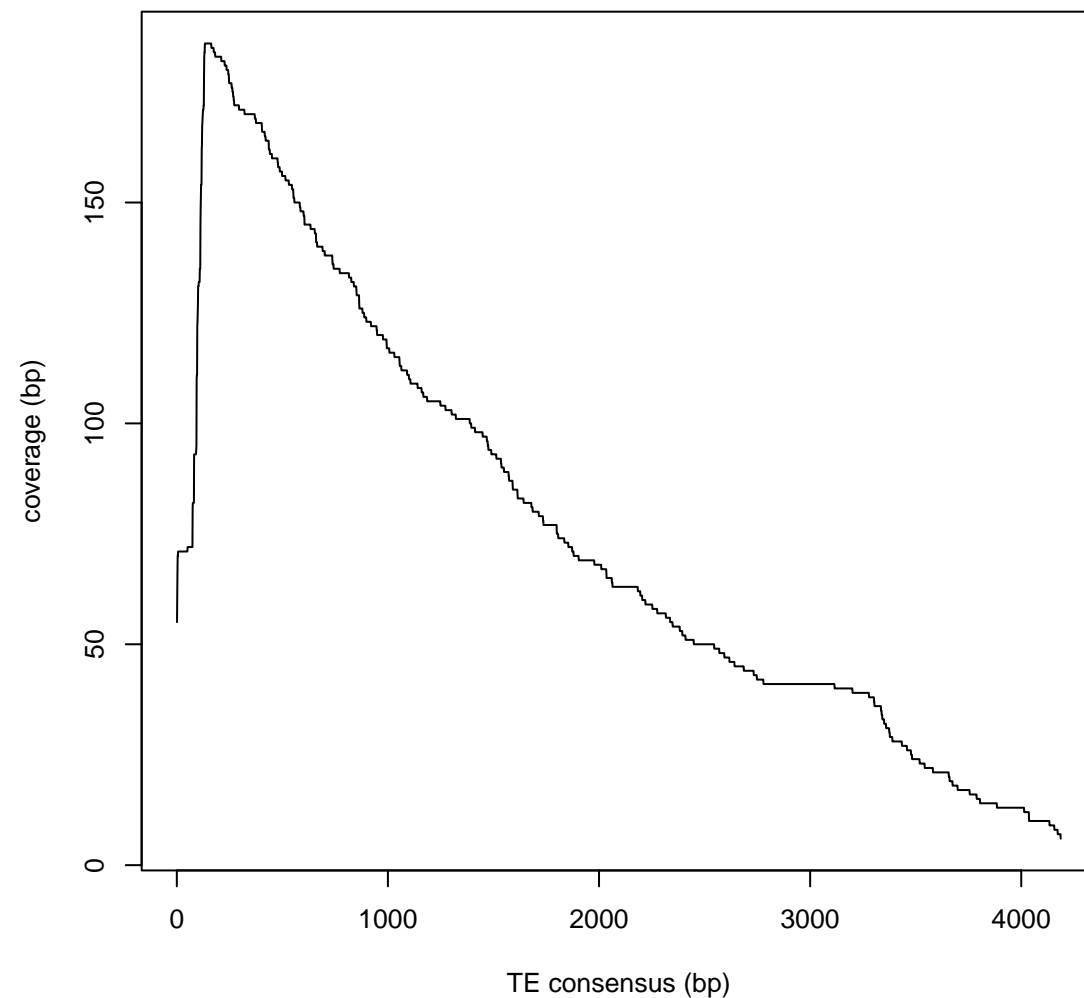


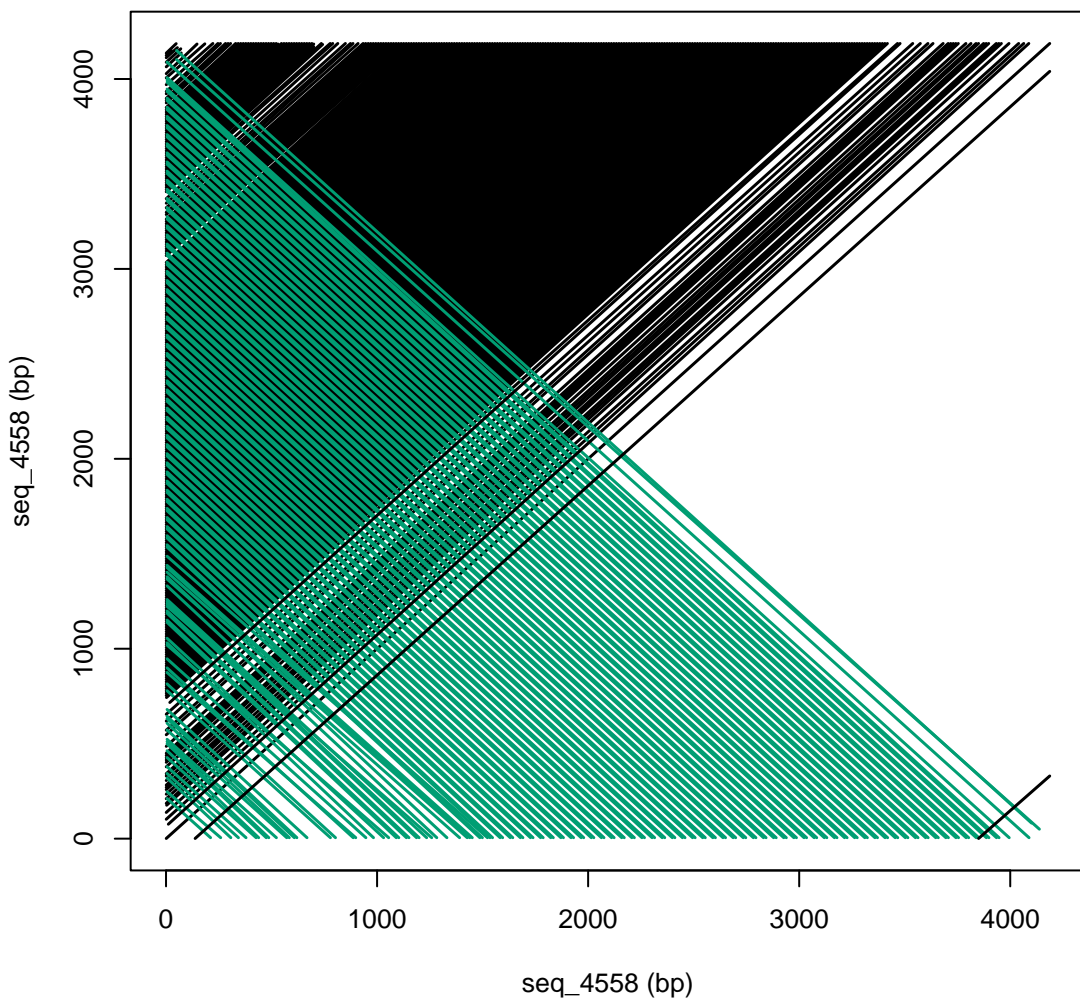
TE: seq\_4558  
consensus size: 4187bp; fragments: 186; full length: 14 ( $\geq 3768.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

