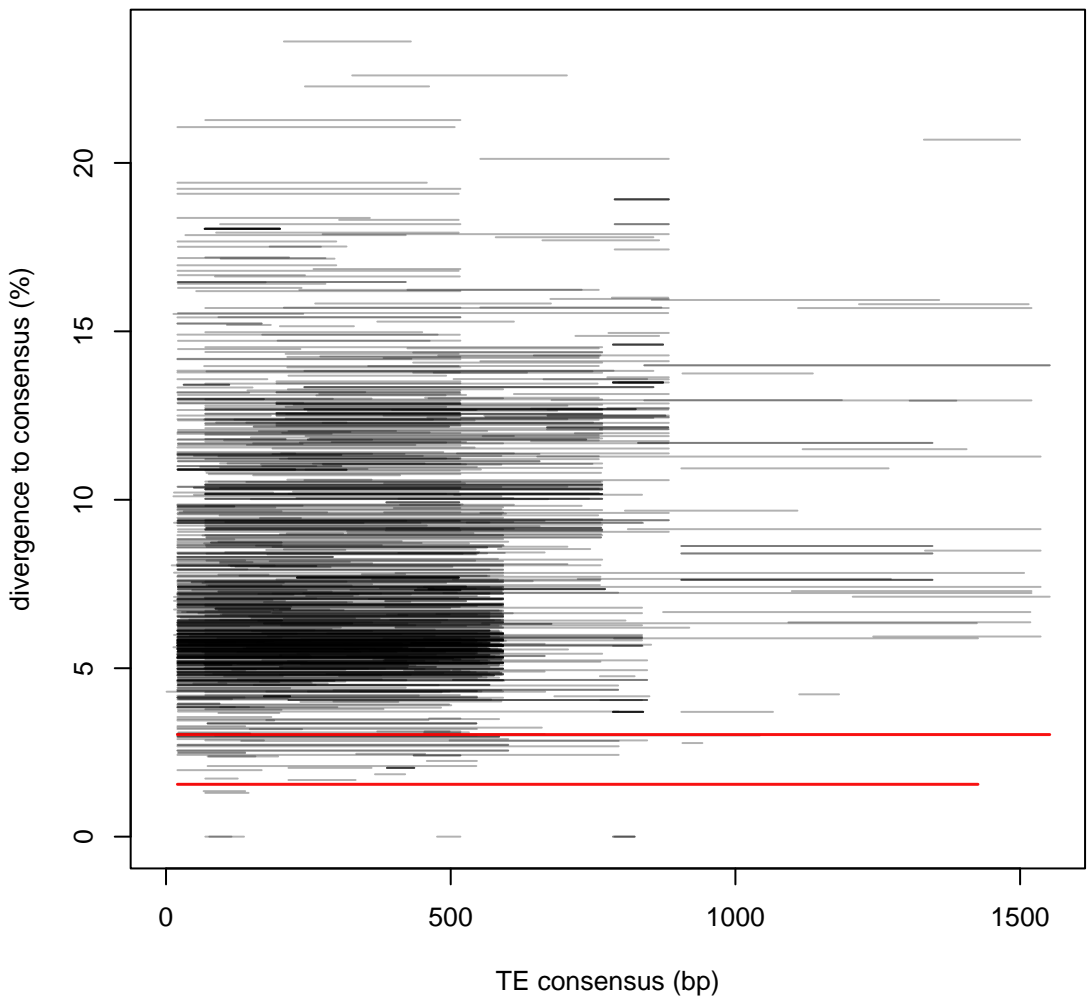
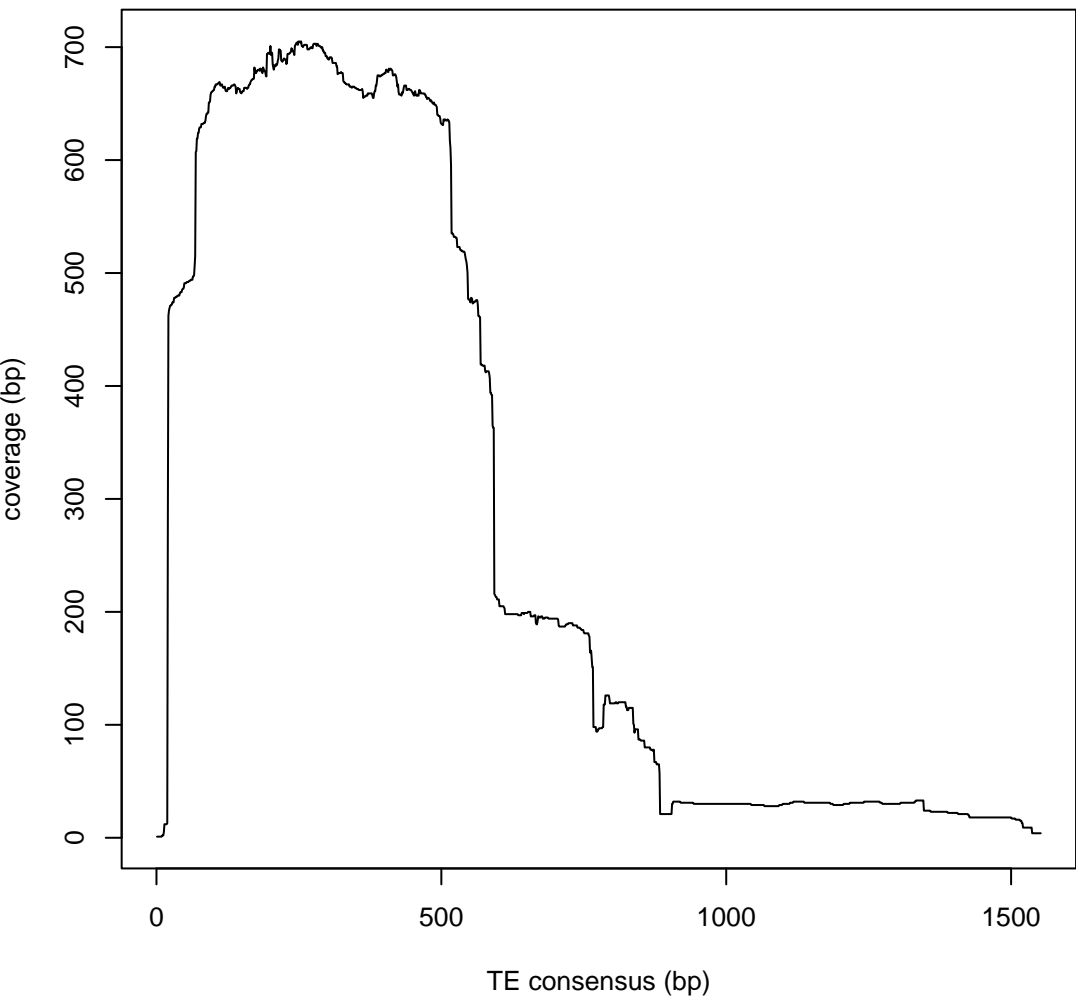


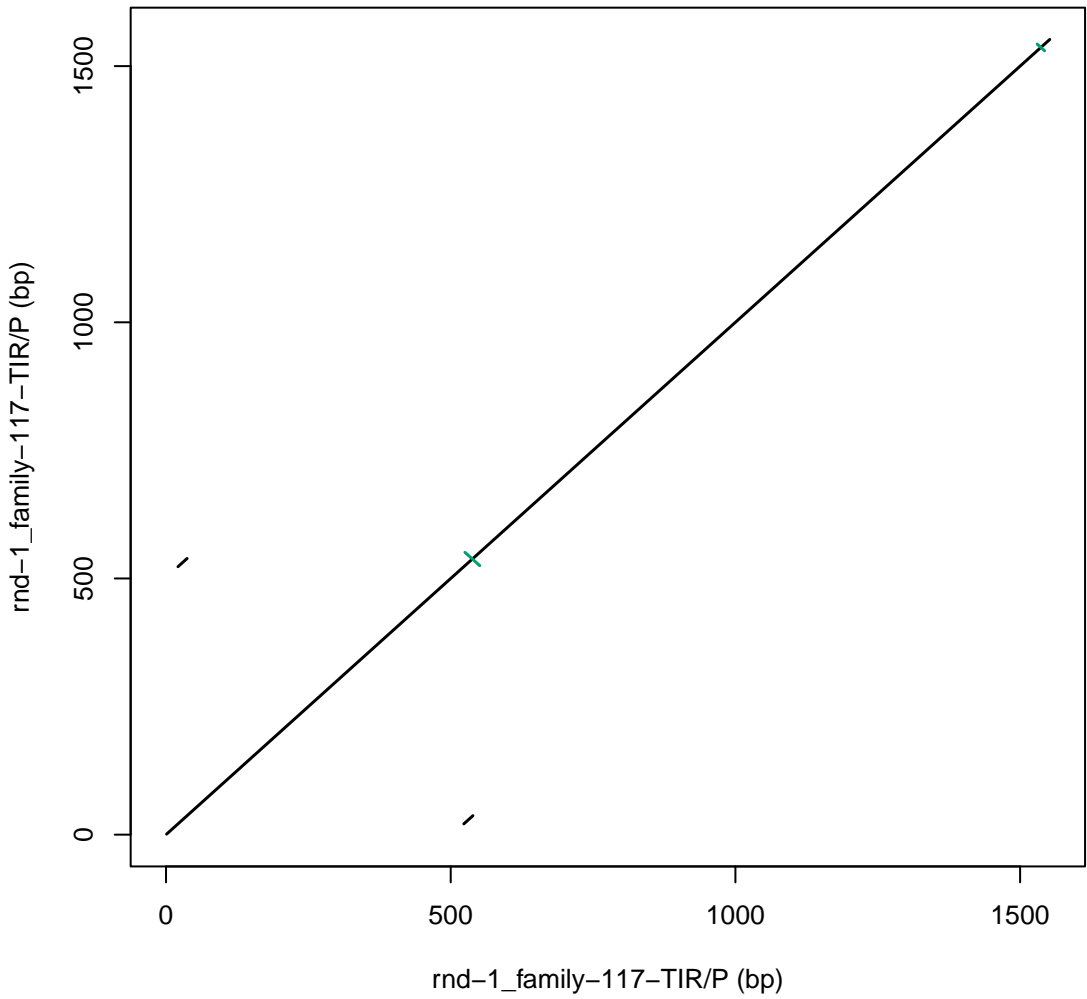
TE: rnd-1\_family-117-TIR/P  
consensus size: 1552bp; fragments: 1050; full length: 2 ( $\geq 1396.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

