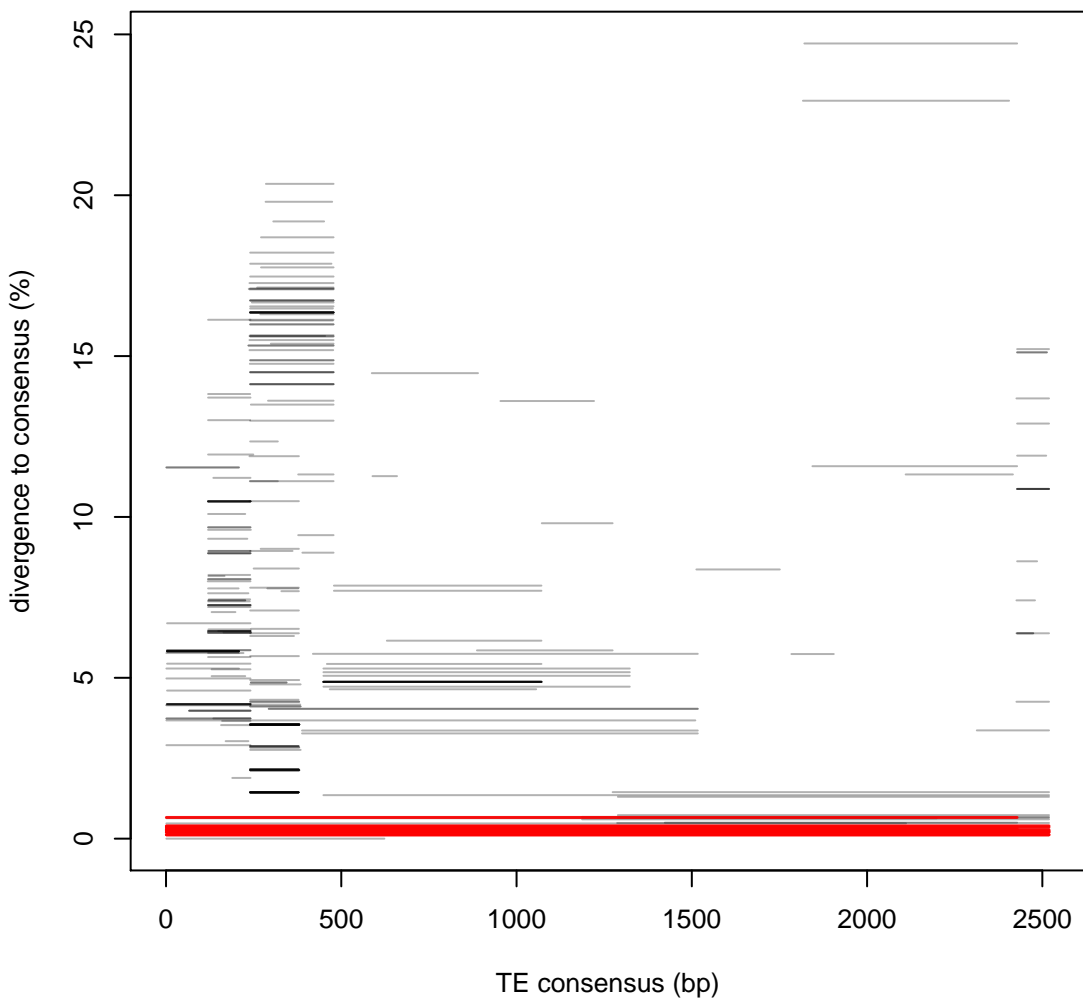
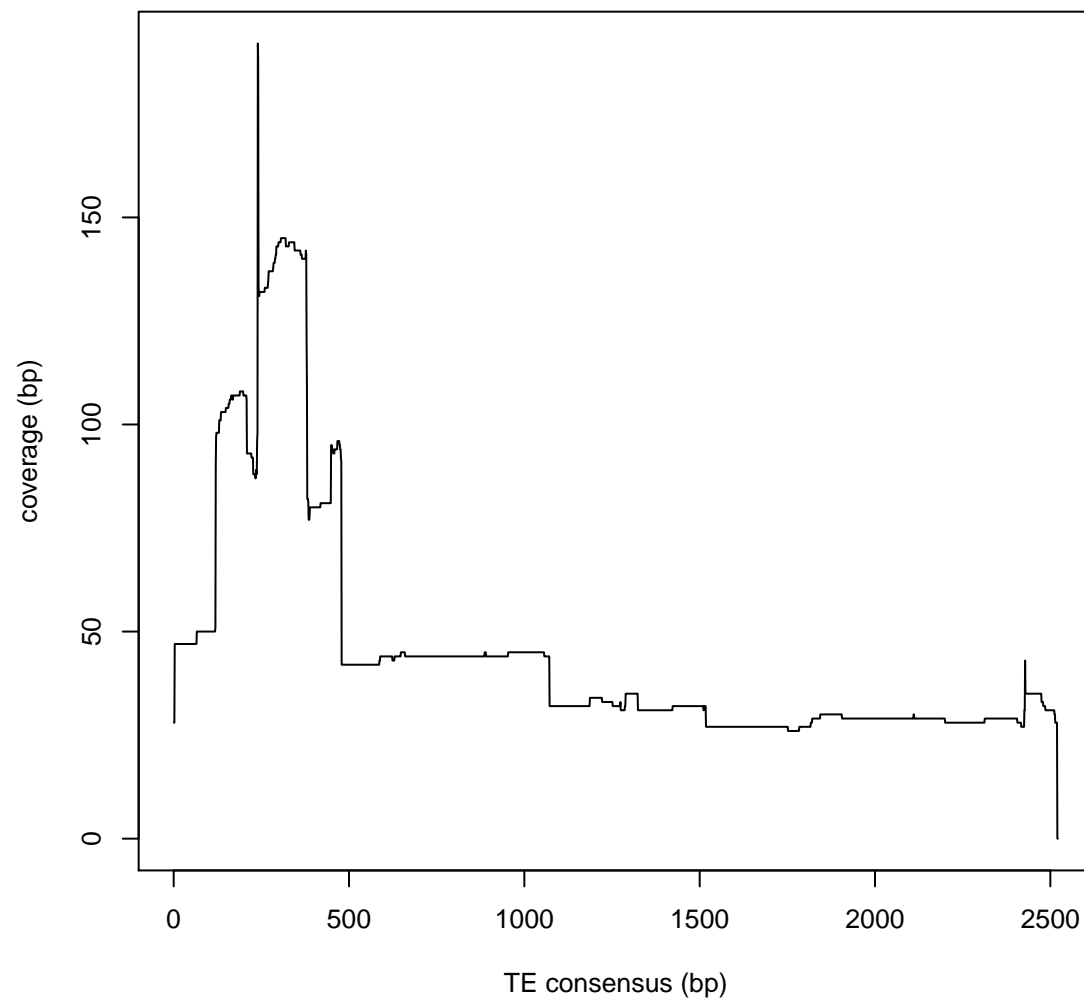


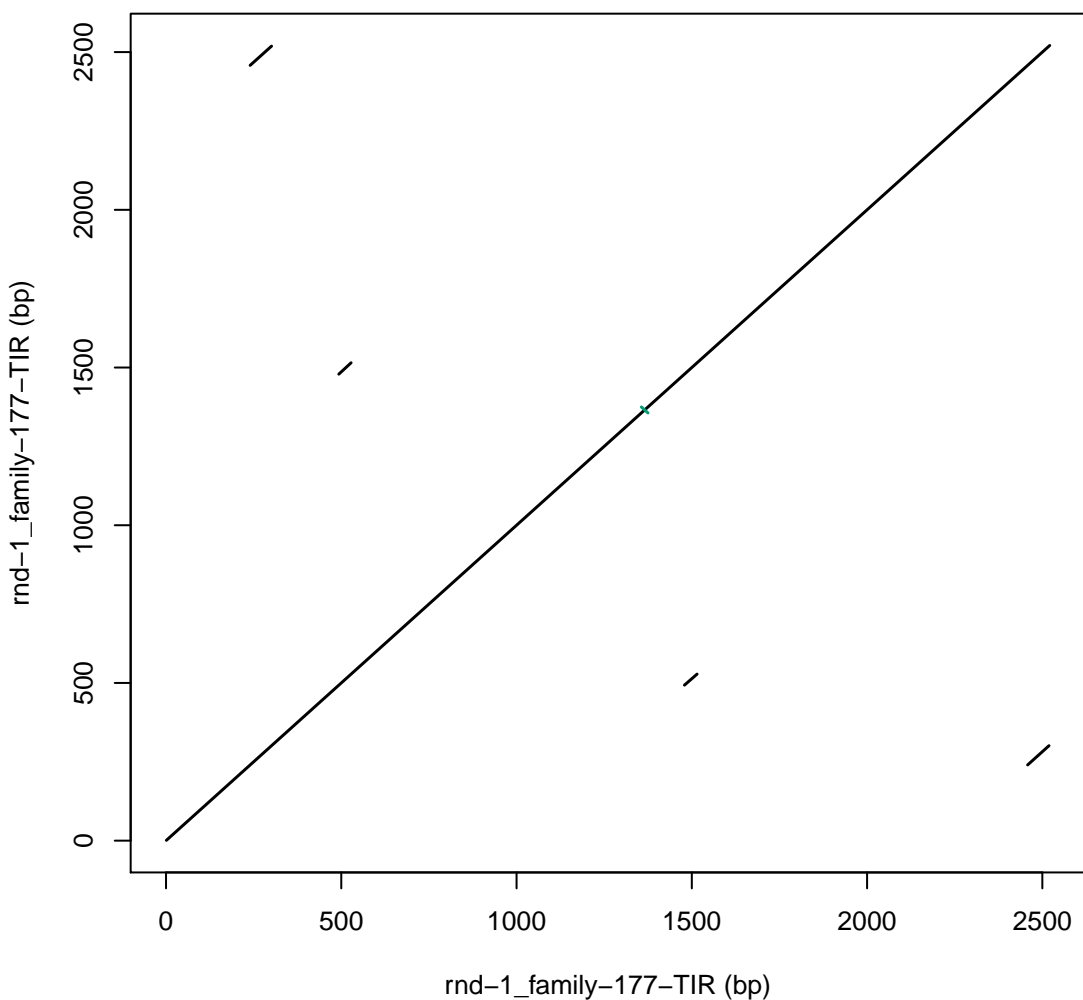
TE: rnd-1\_family-177-TIR  
consensus size: 2521bp; fragments: 300; full length: 14 ( $\geq 2268.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

