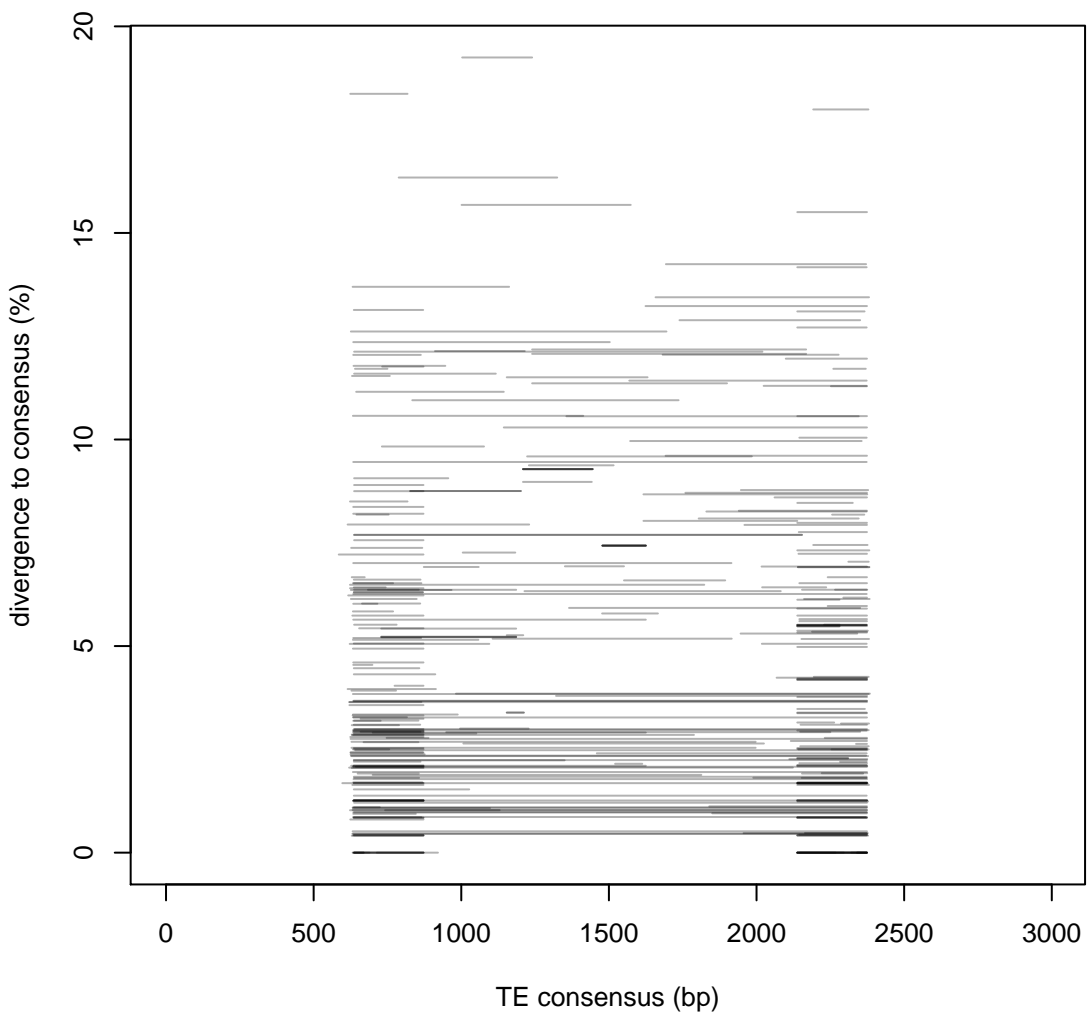
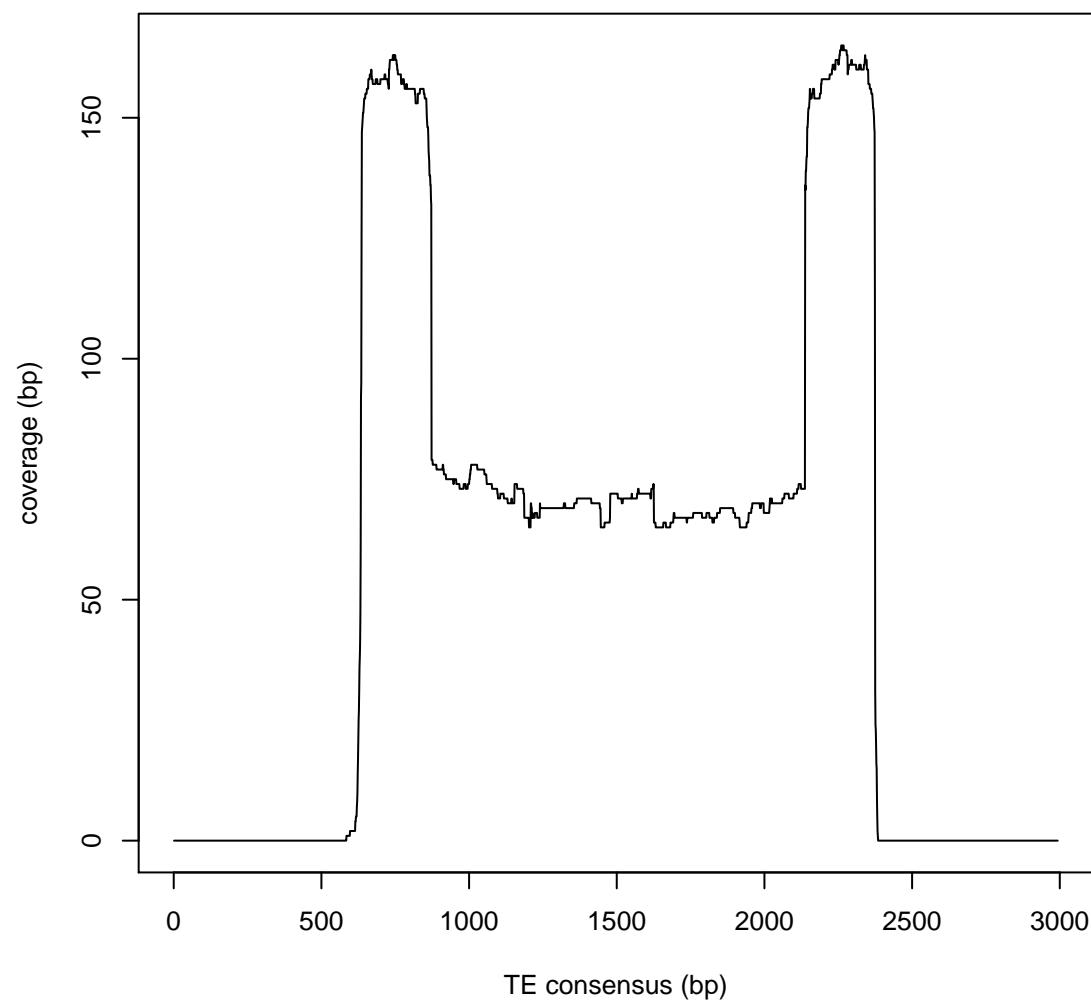


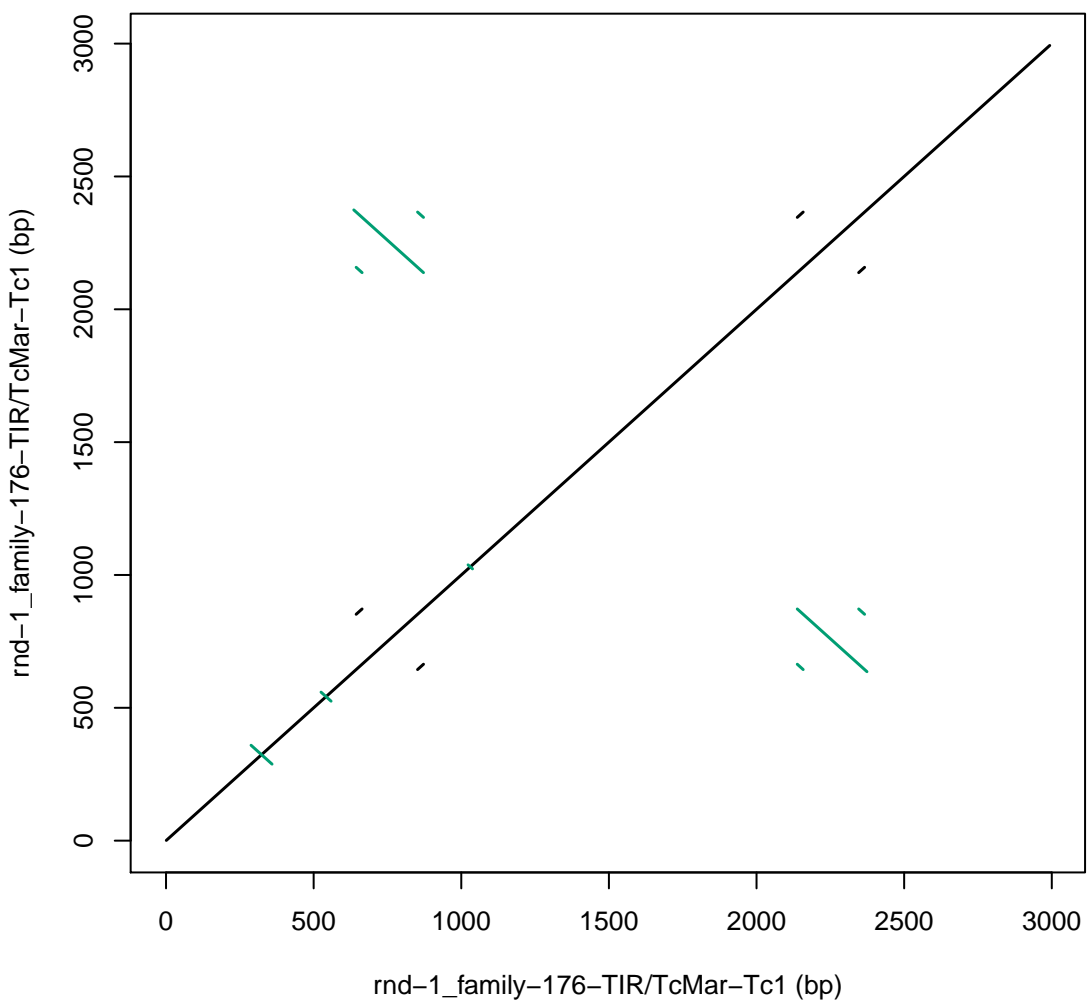
TE: rnd-1\_family-176-TIR/TcMar-Tc1  
consensus size: 2993bp; fragments: 354; full length: 0 (>=2693.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

