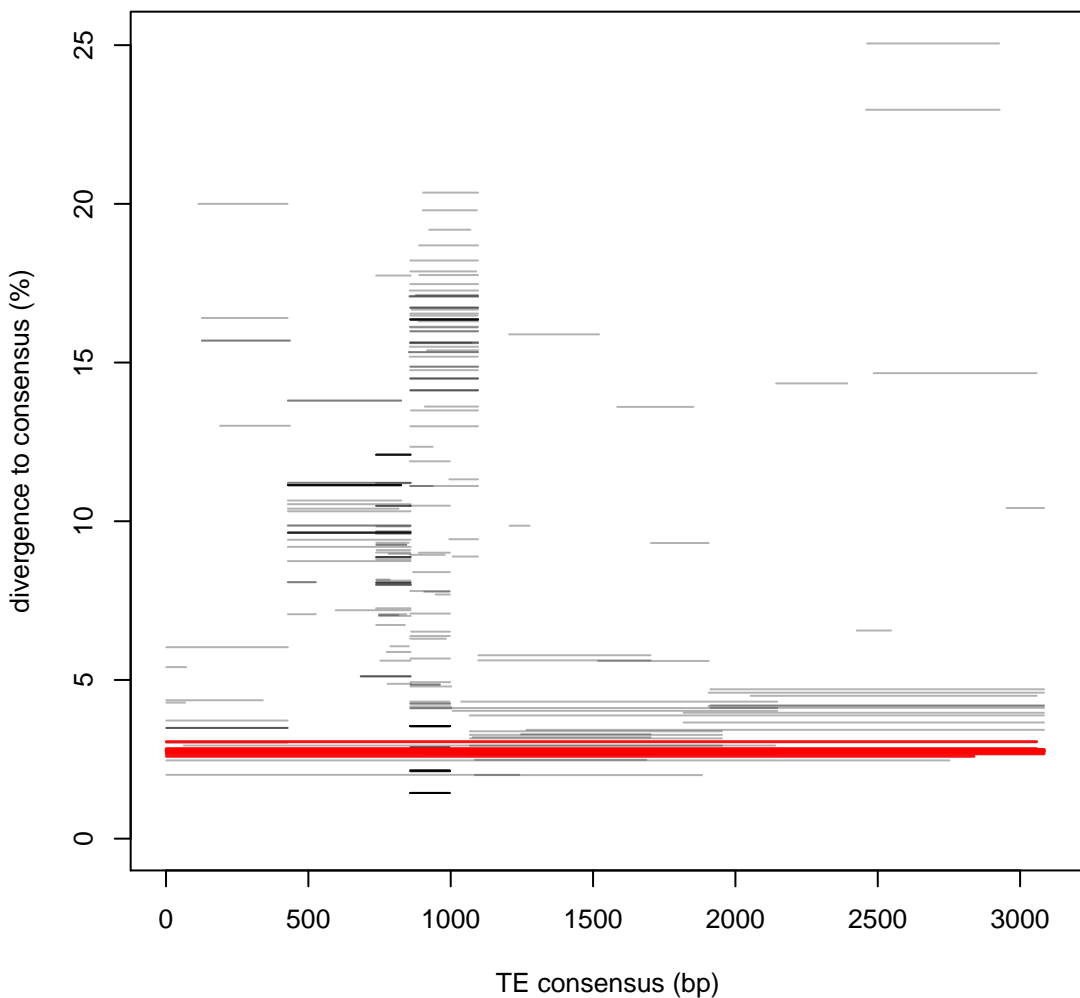
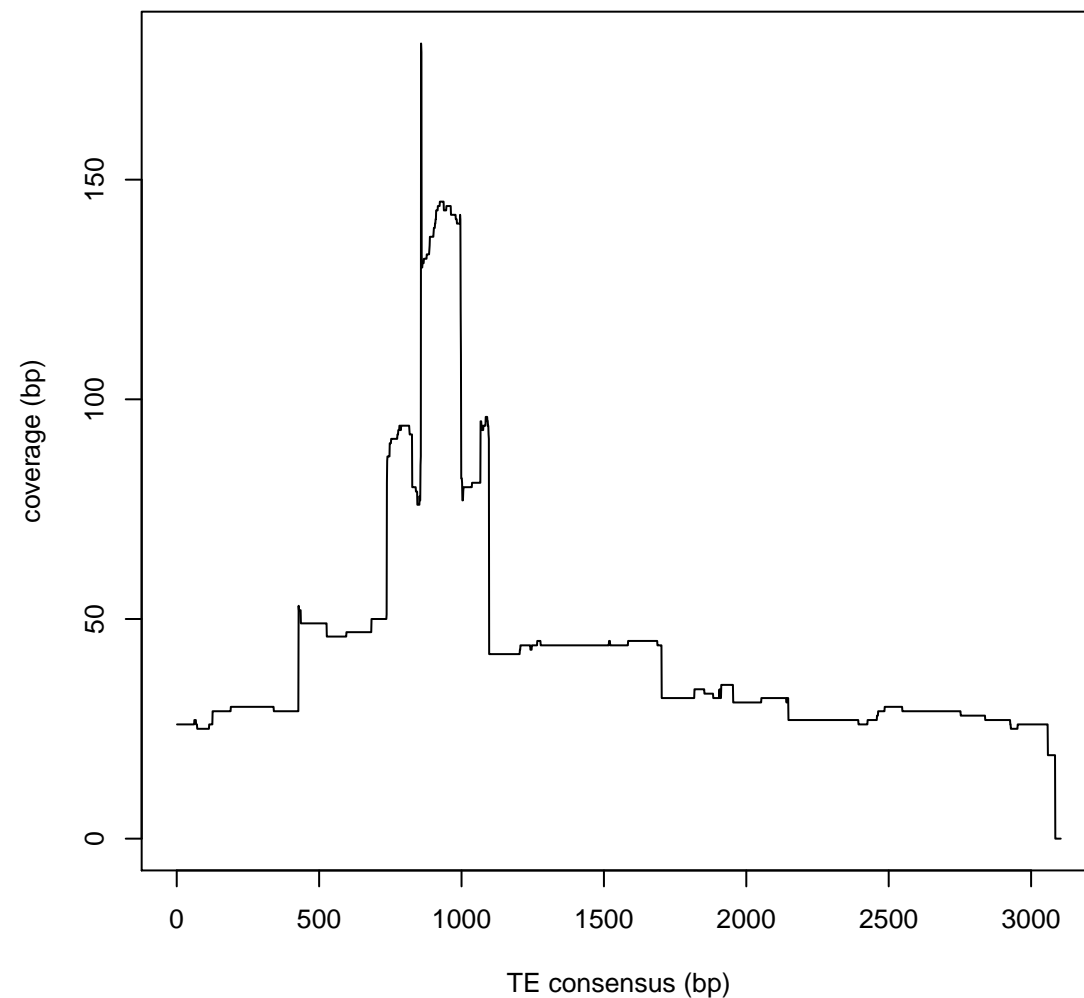


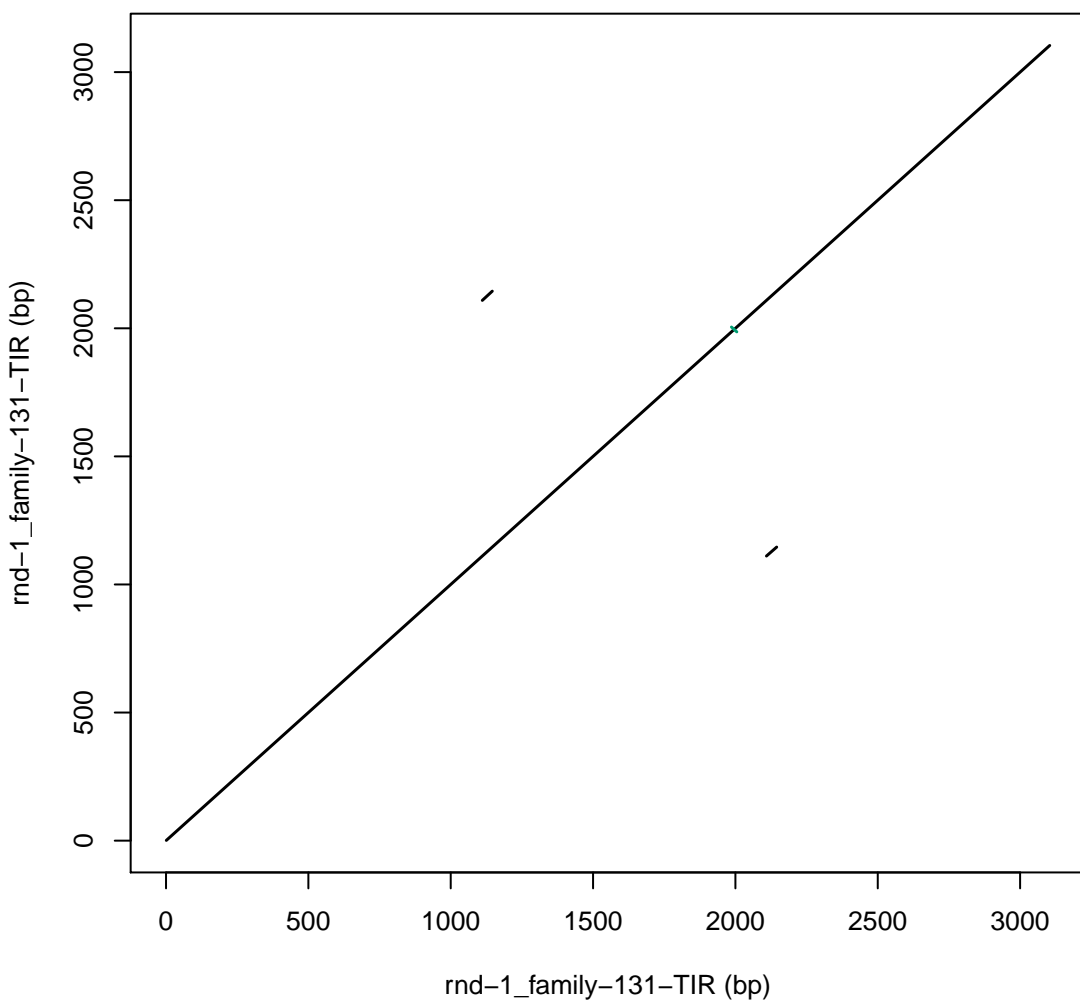
TE: rnd-1_family-131-TIR
consensus size: 3104bp; fragments: 286; full length: 15 (≥ 2793.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

