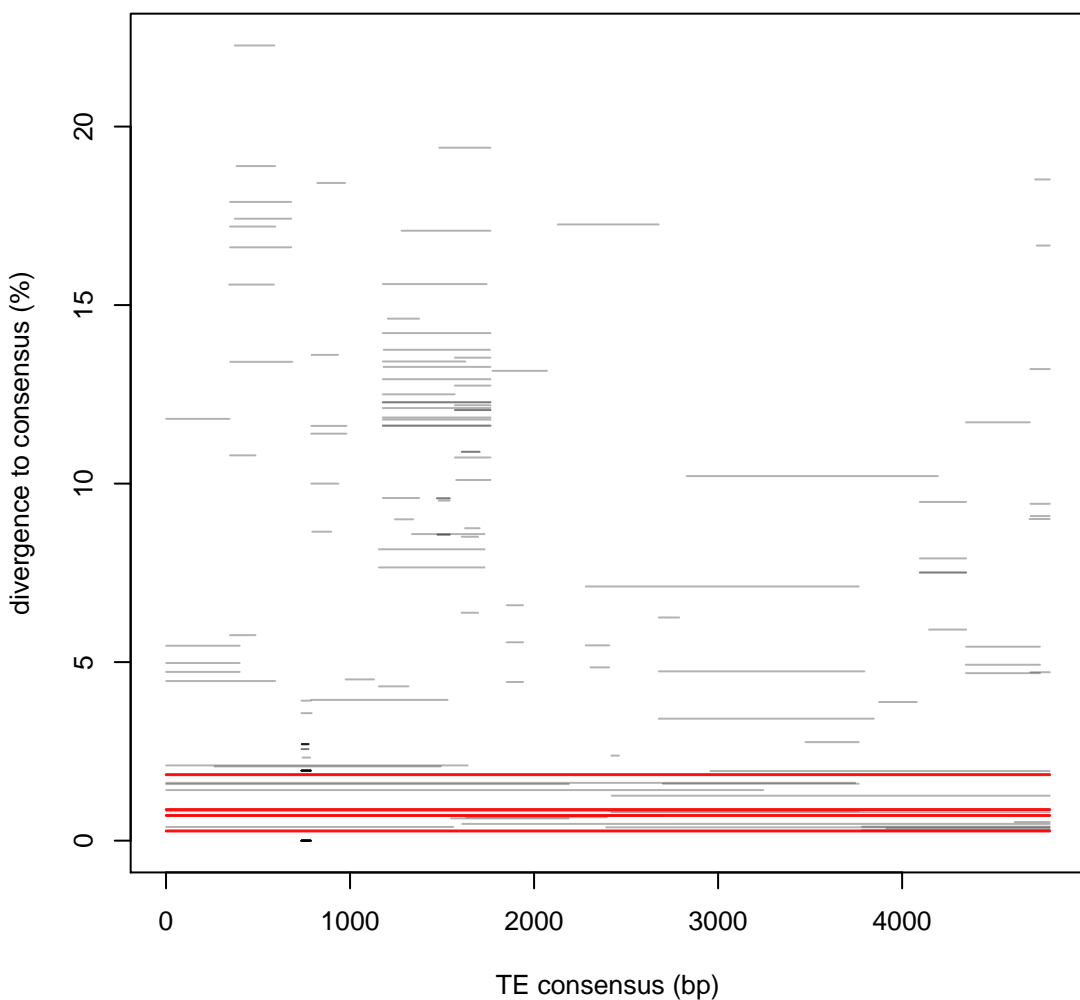
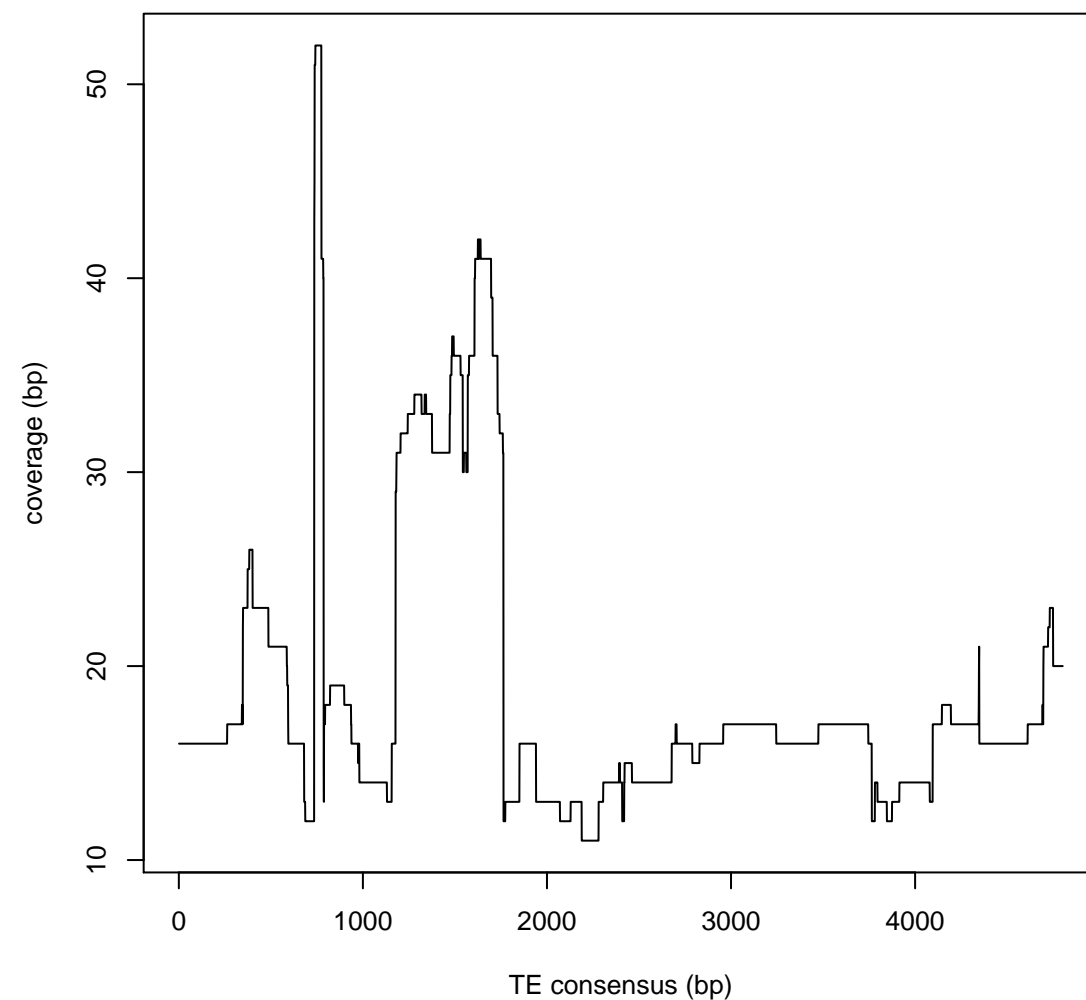


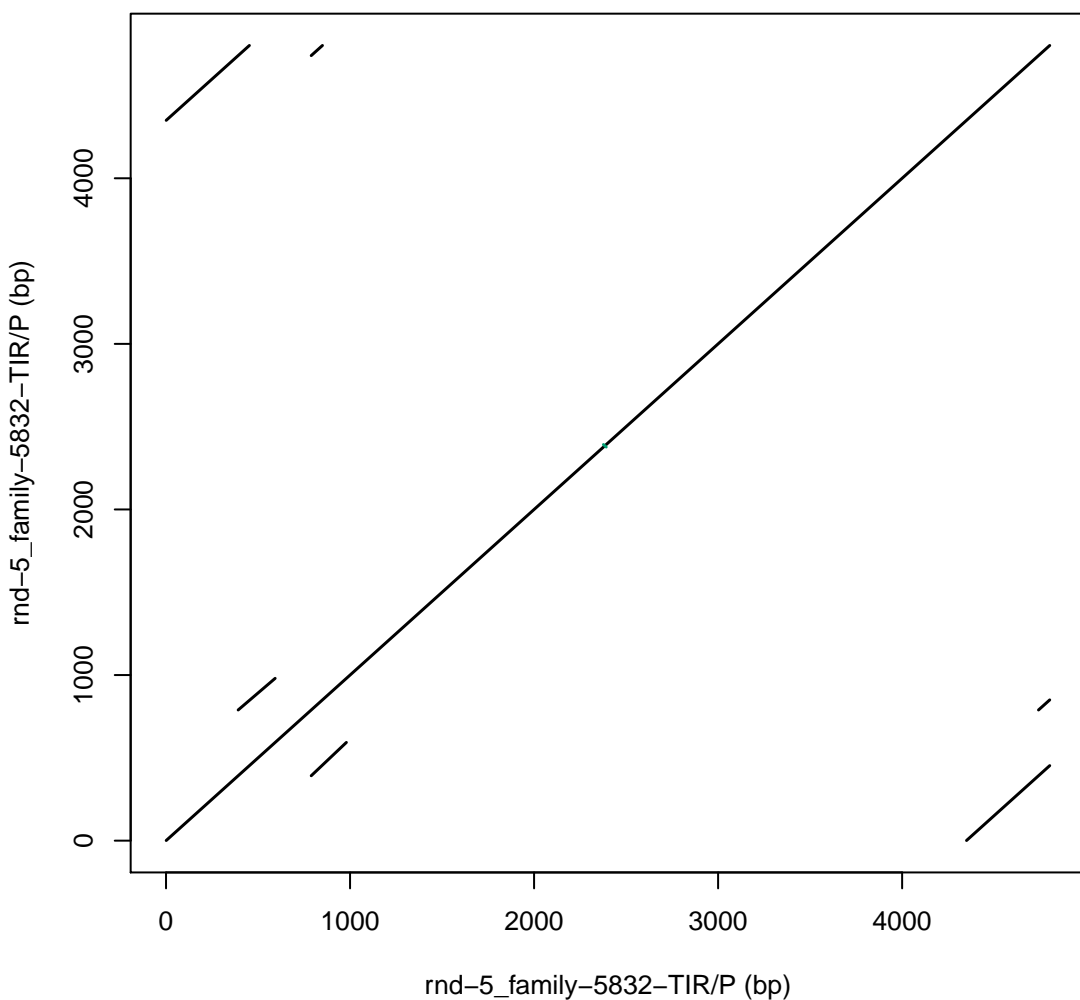
TE: rnd-5_family-5832-TIR/P
consensus size: 4802bp; fragments: 158; full length: 4 (≥ 4321.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

