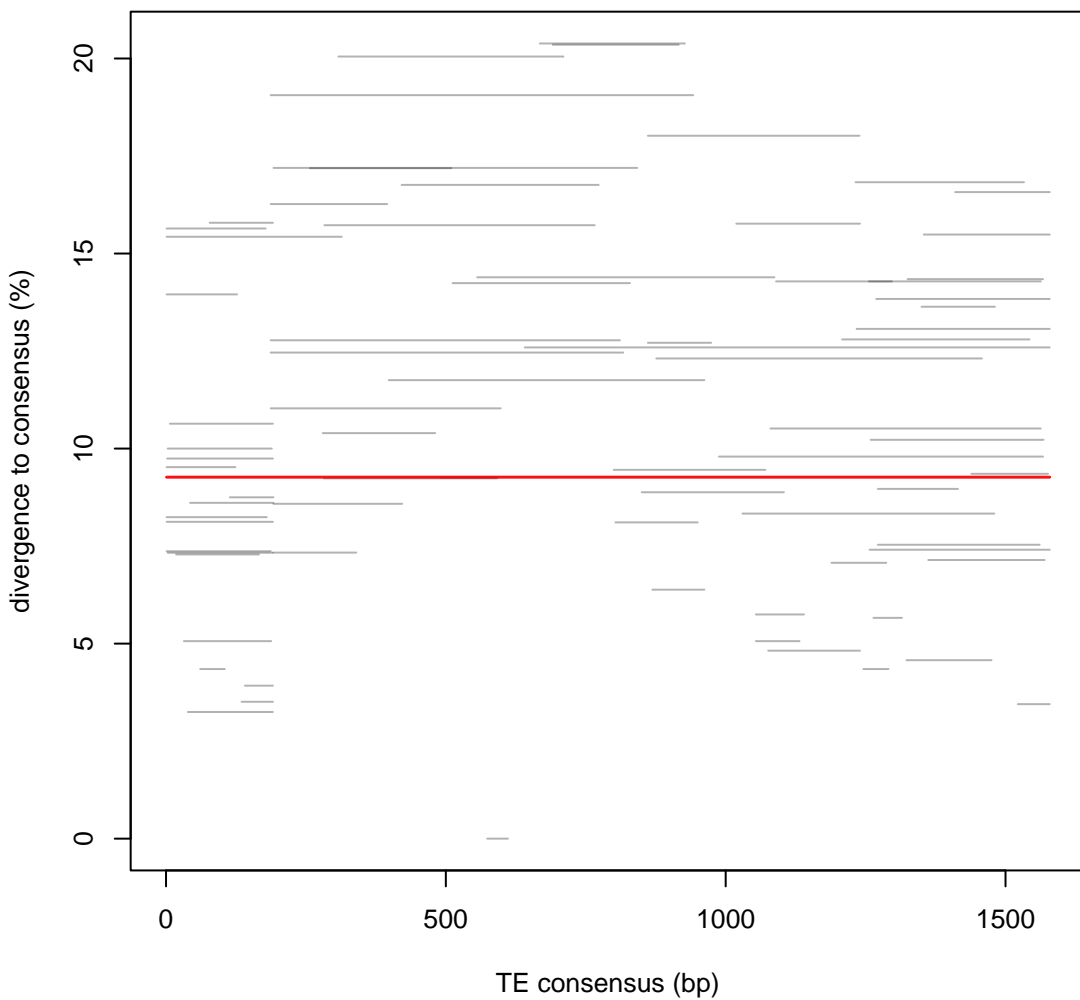
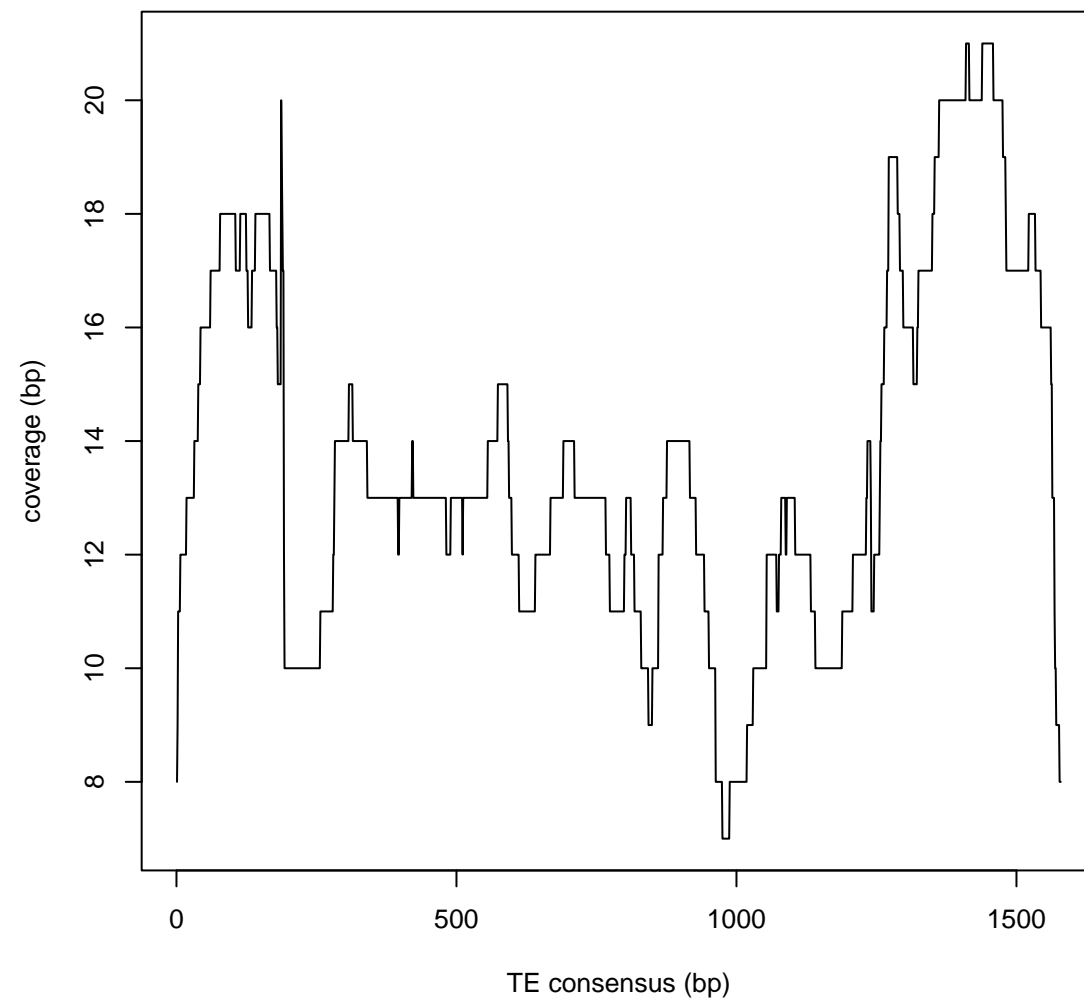


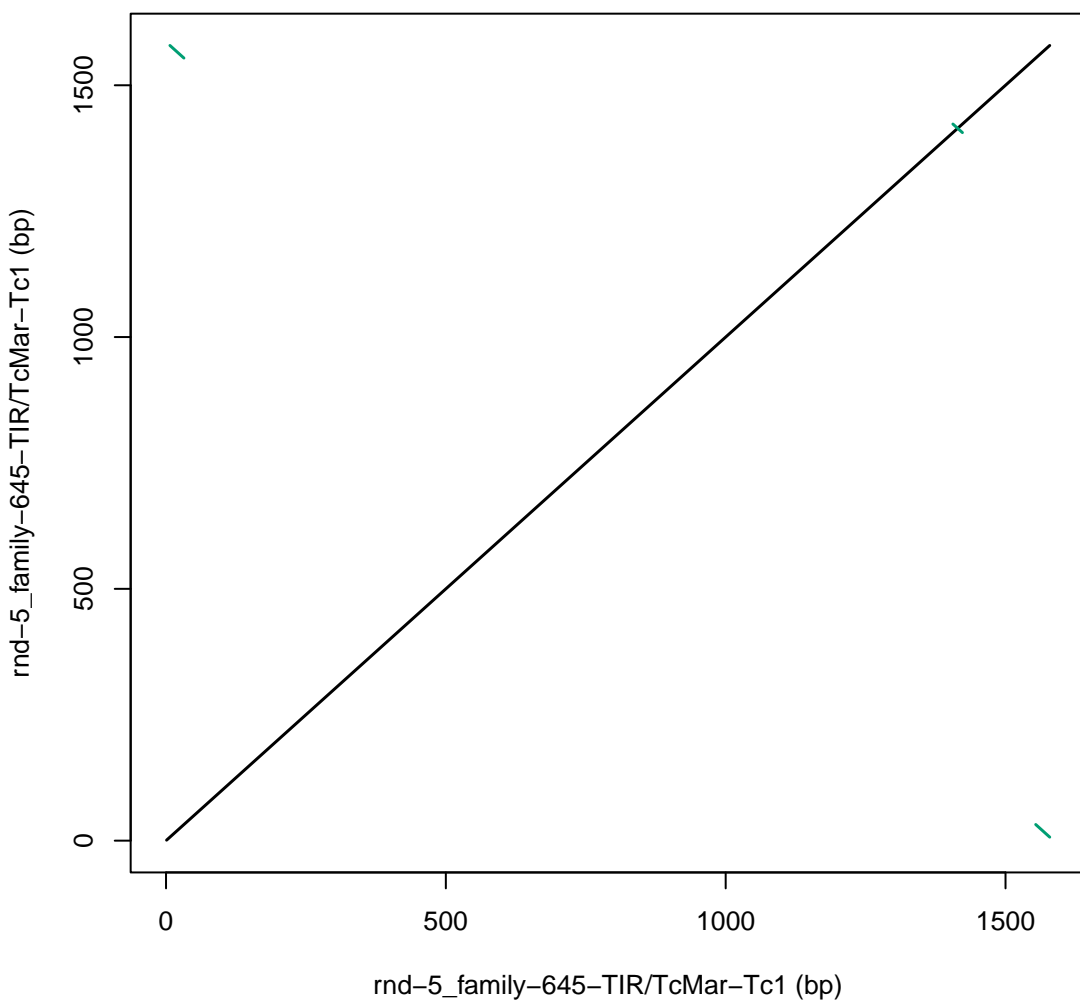
TE: rnd-5_family-645-TIR/TcMar-Tc1
consensus size: 1579bp; fragments: 78; full length: 1 (≥ 1421.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

