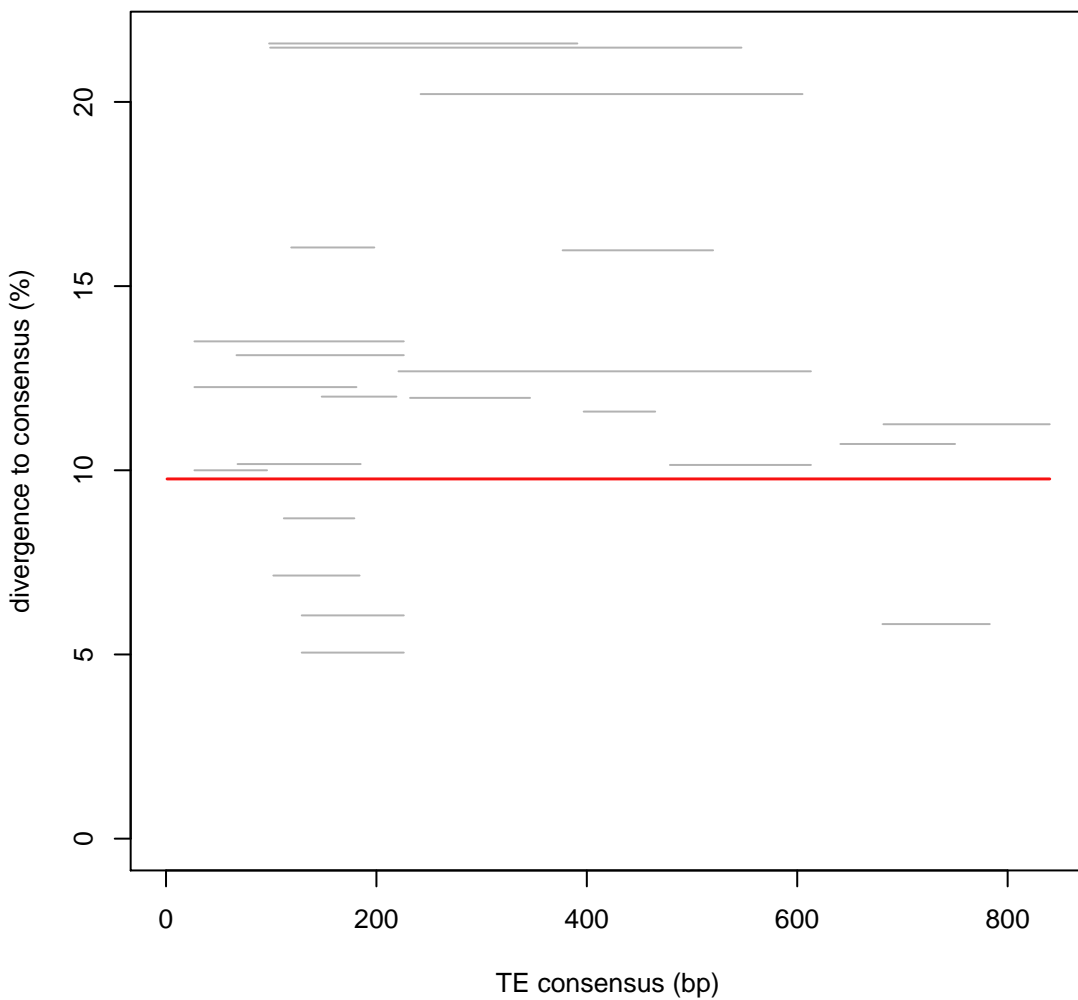
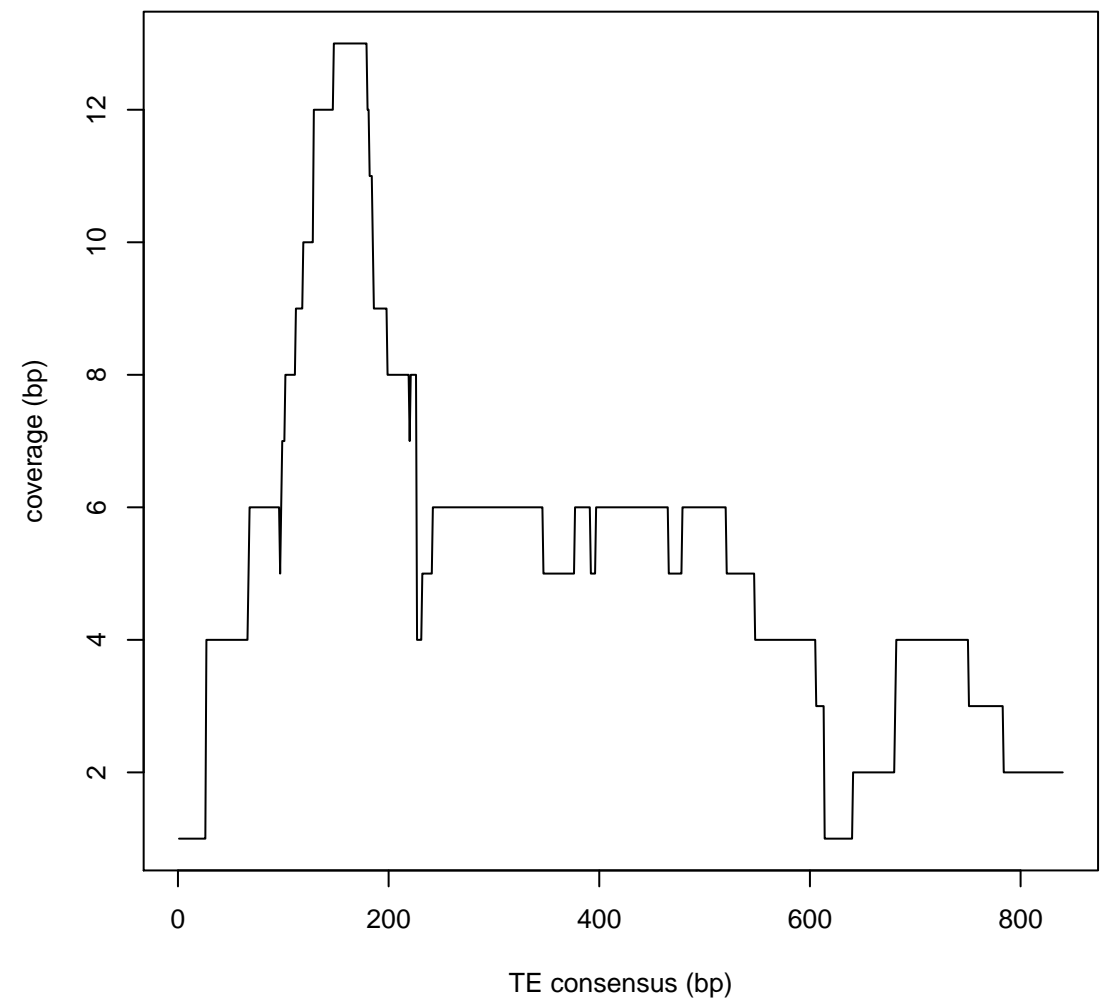


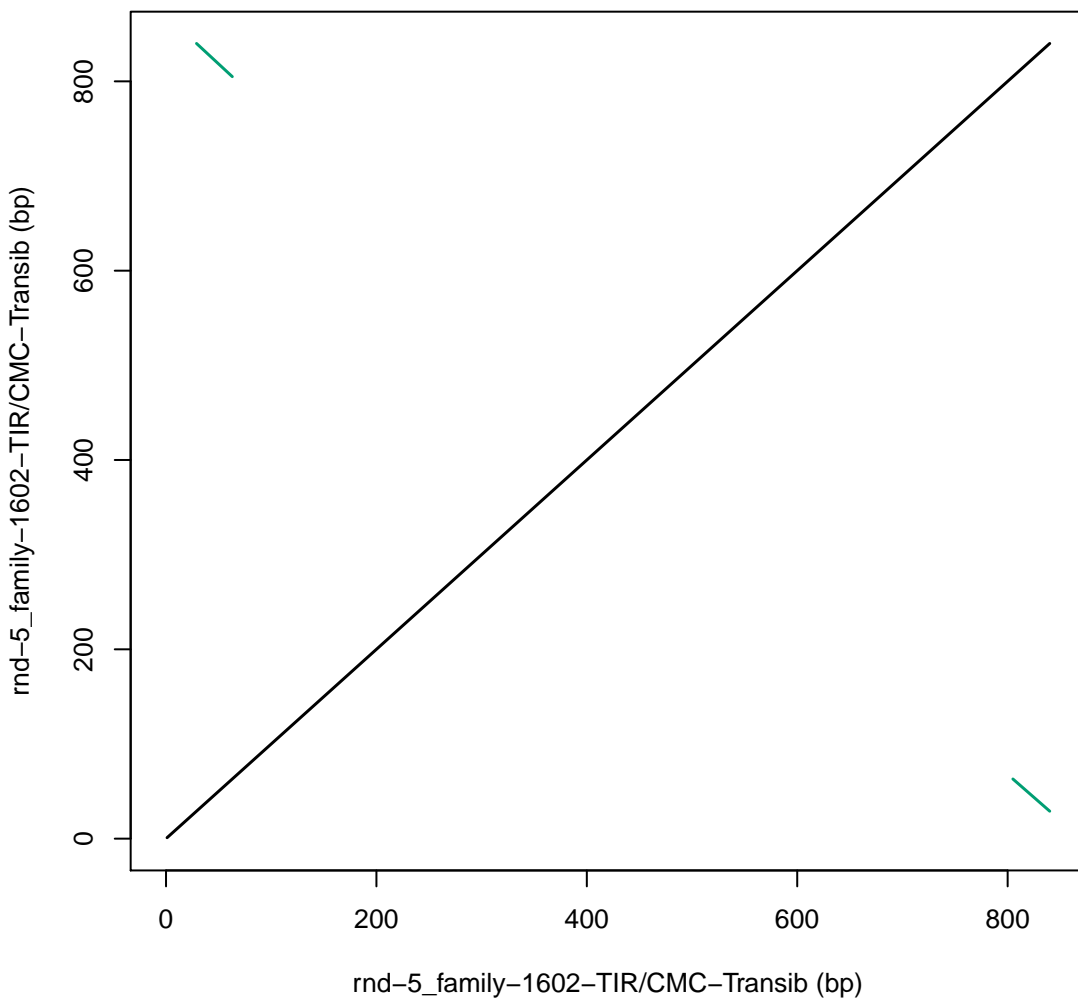
TE: rnd-5_family-1602-TIR/CMC-Transib
consensus size: 840bp; fragments: 23; full length: 1 (>=756bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

