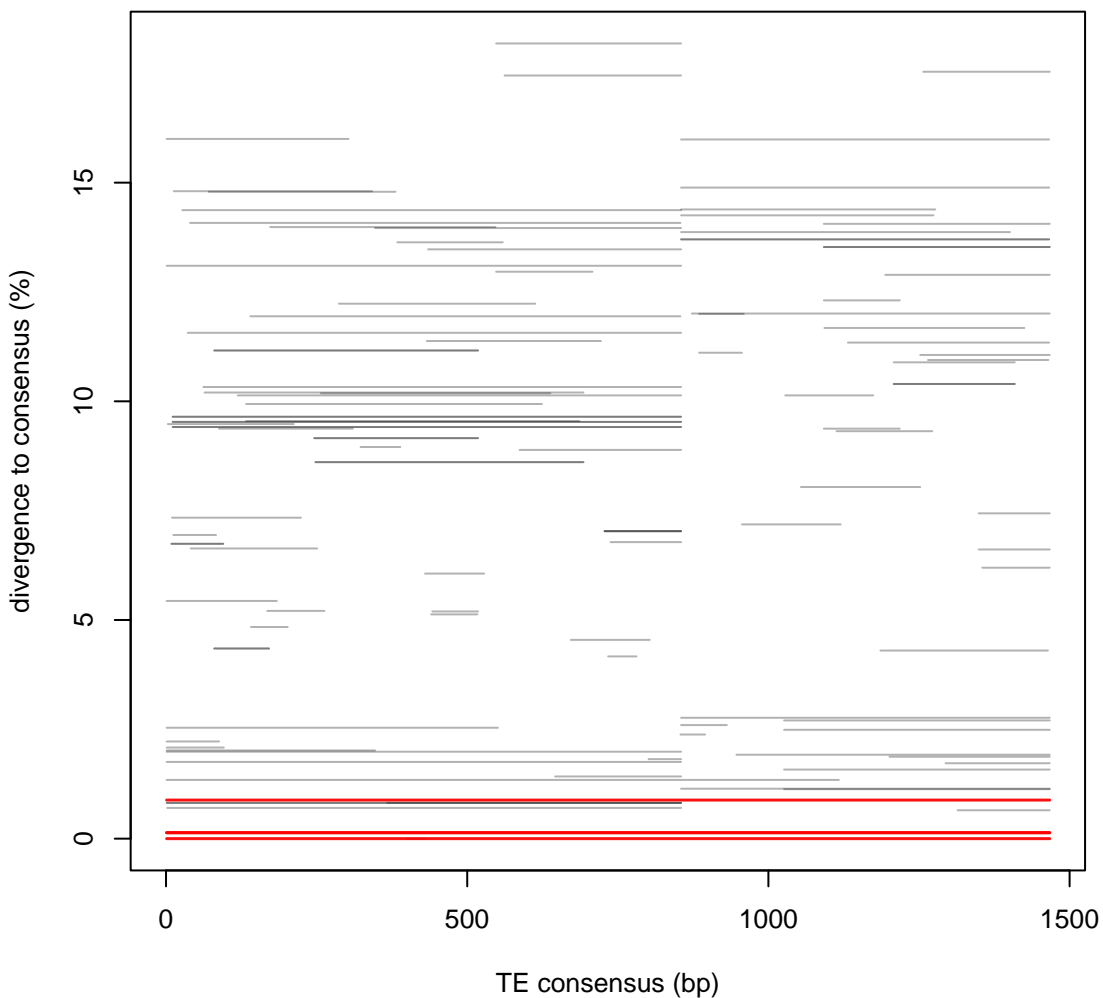
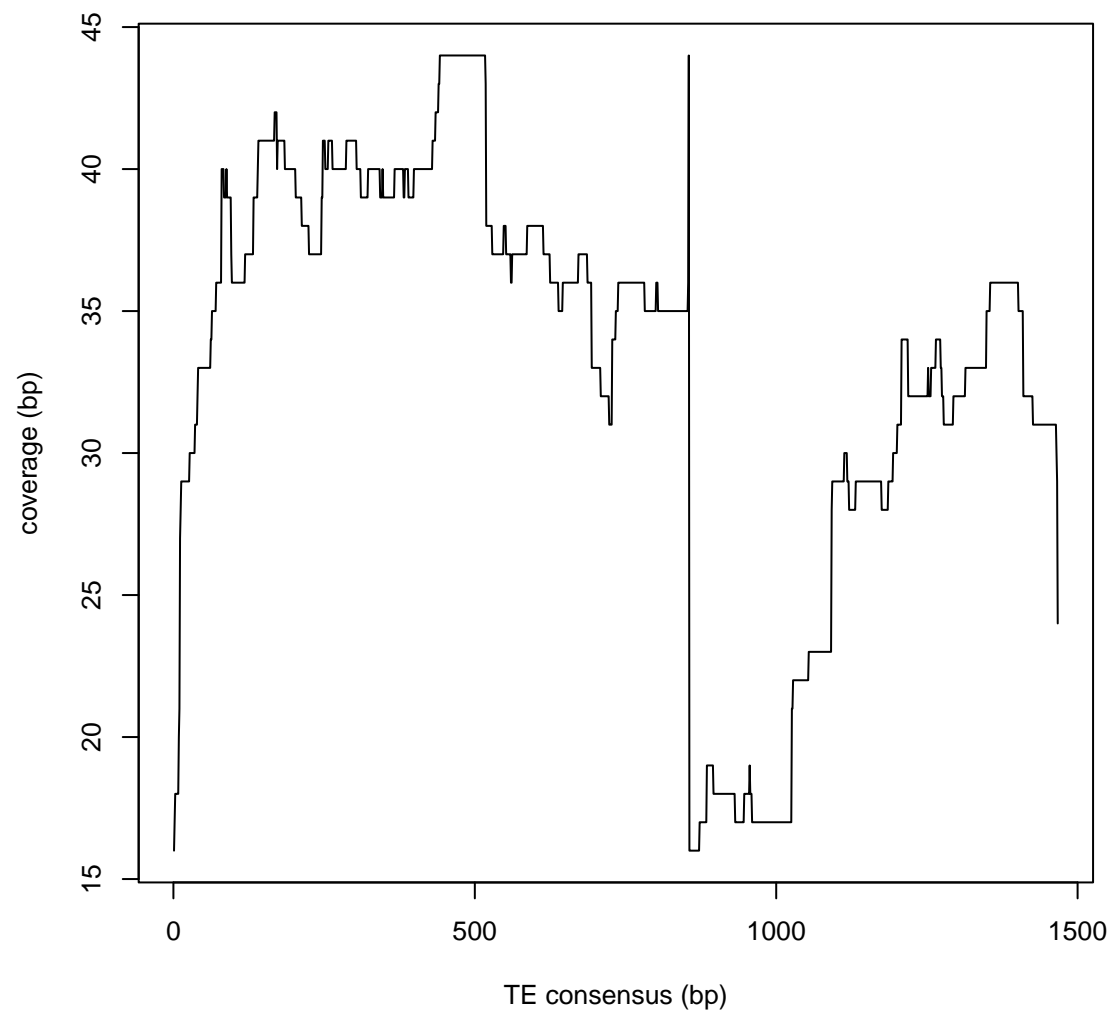


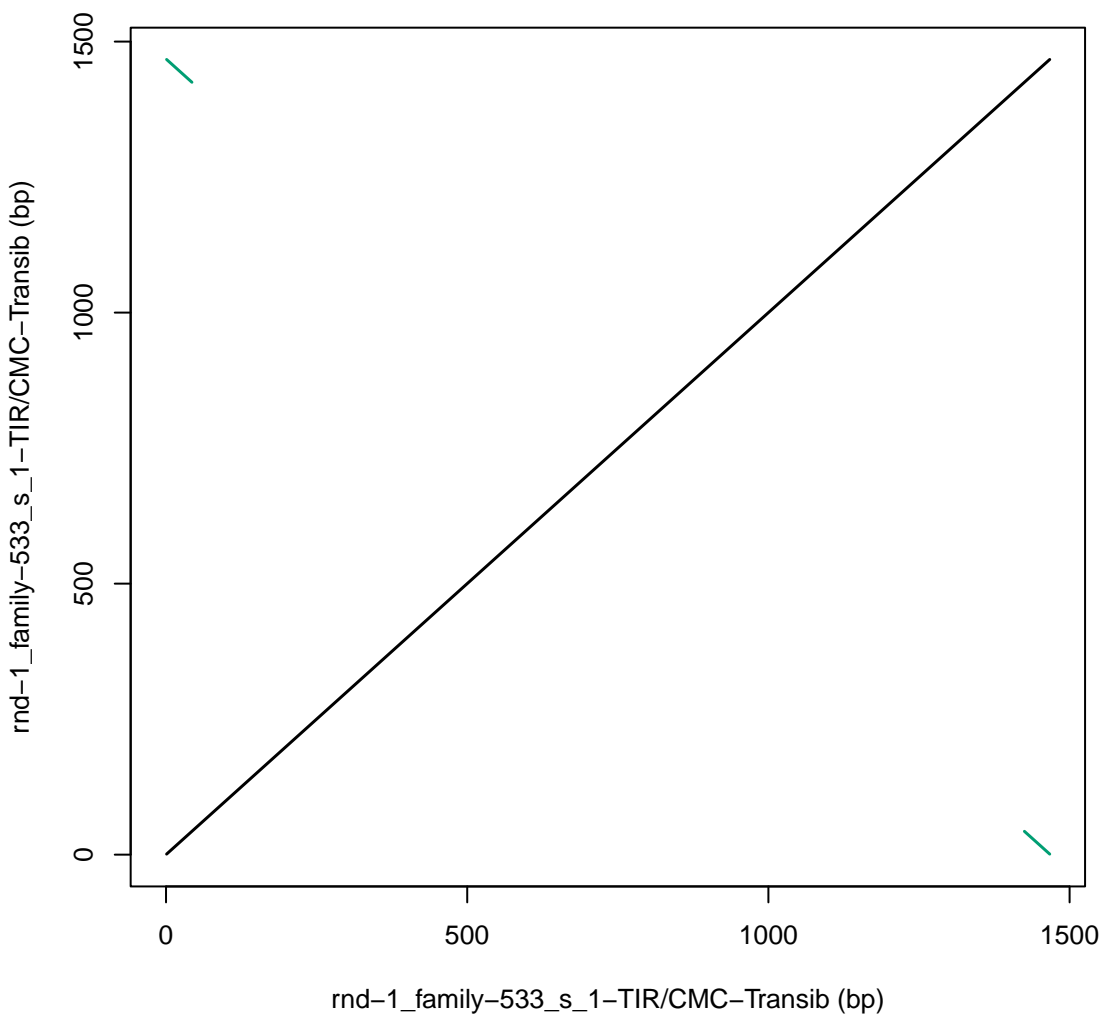
TE: rnd-1_family-533_s_1-TIR/CMC-Transib
consensus size: 1467bp; fragments: 120; full length: 4 (>=1320.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

