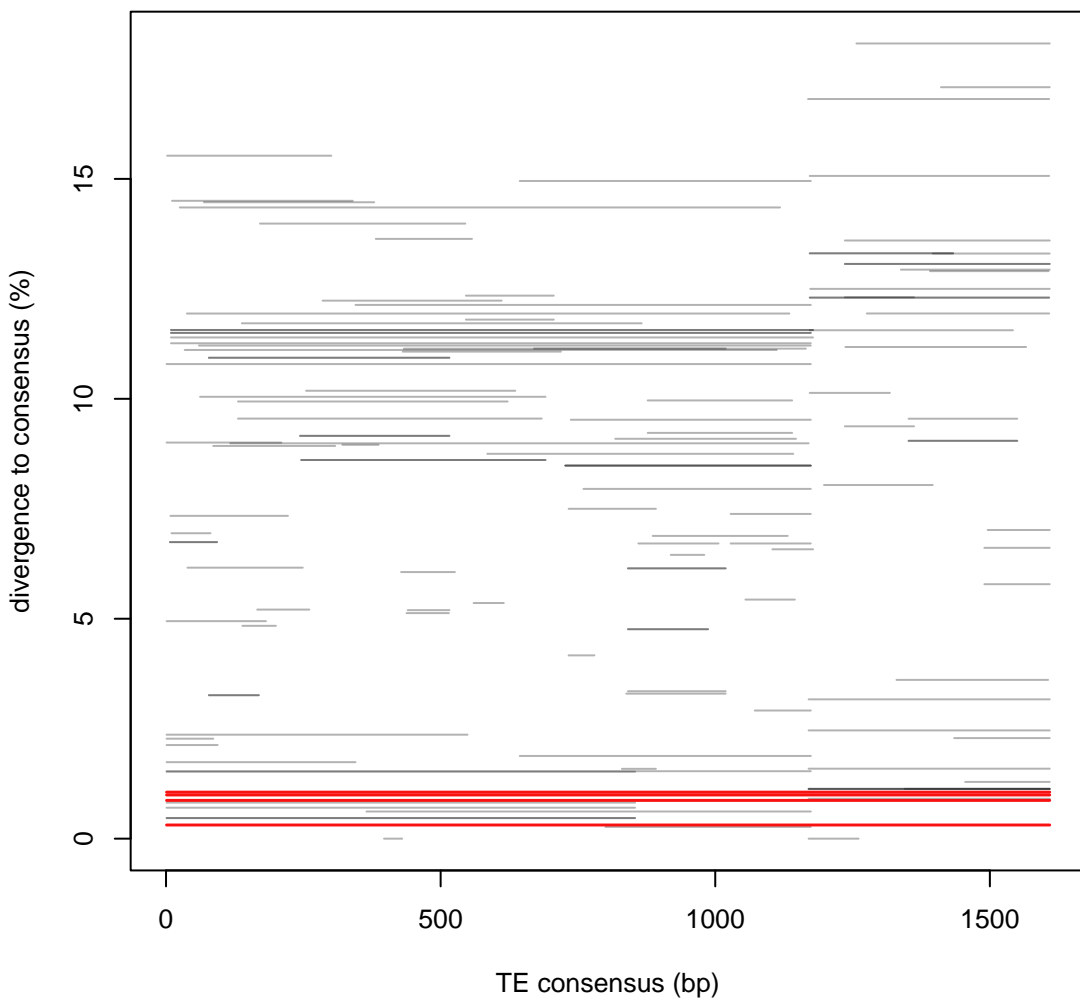
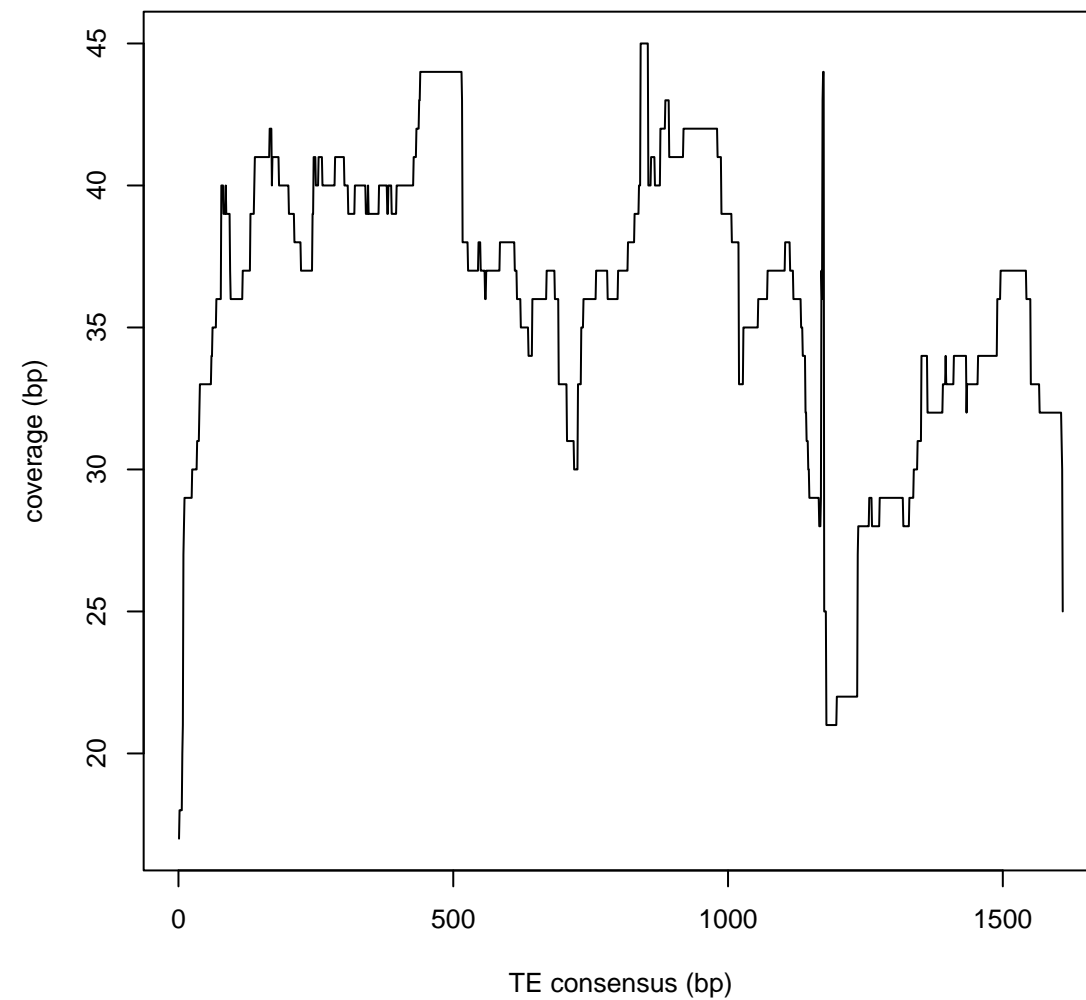


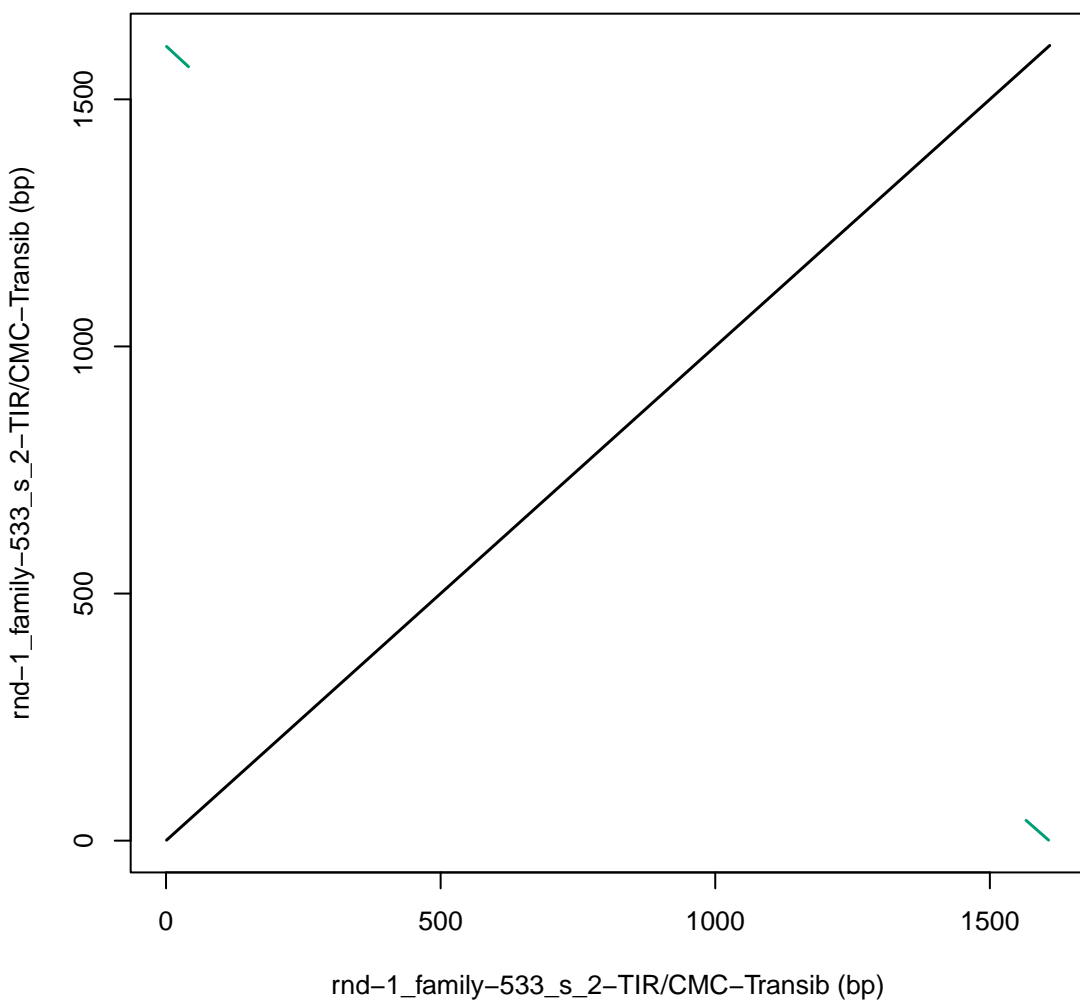
TE: rnd-1_family-533_s_2-TIR/CMC-Transib
consensus size: 1609bp; fragments: 137; full length: 4 (≥ 1448.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

