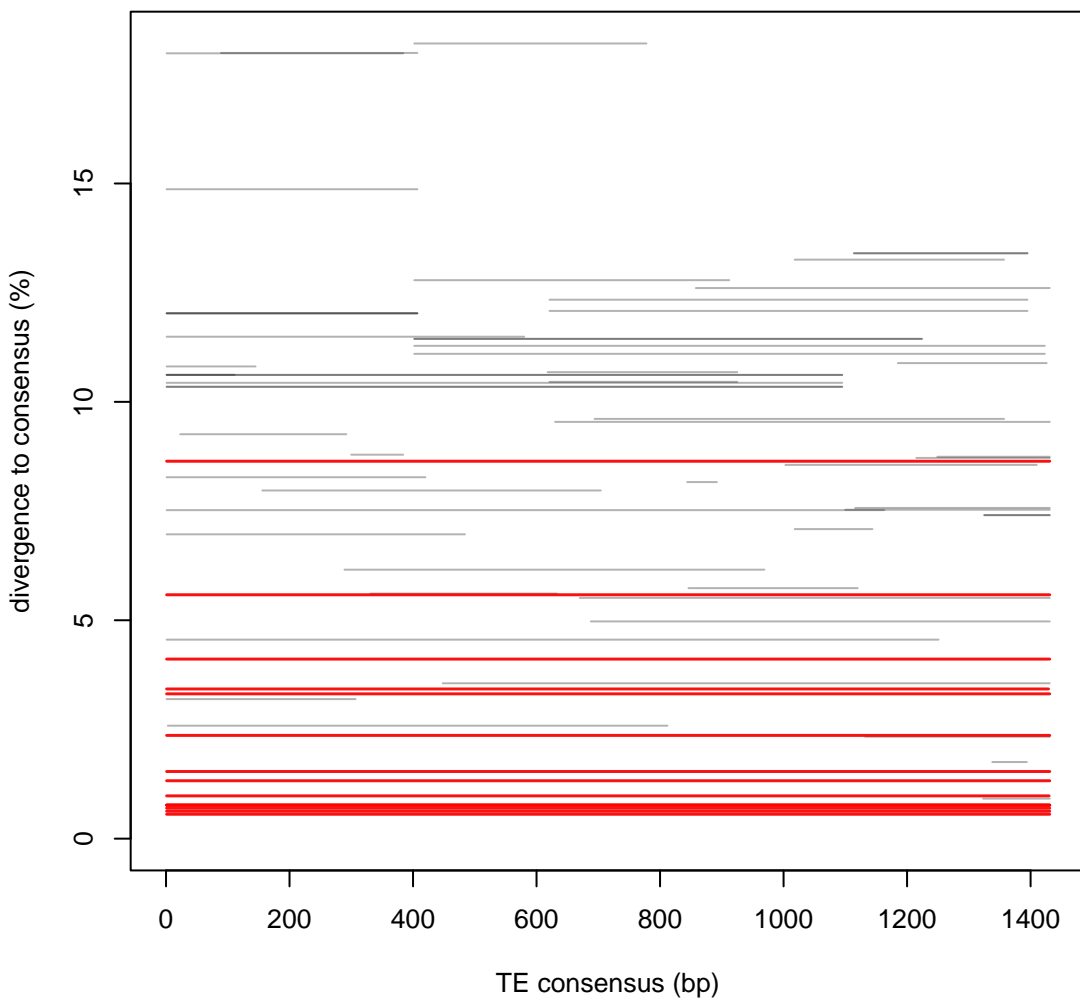
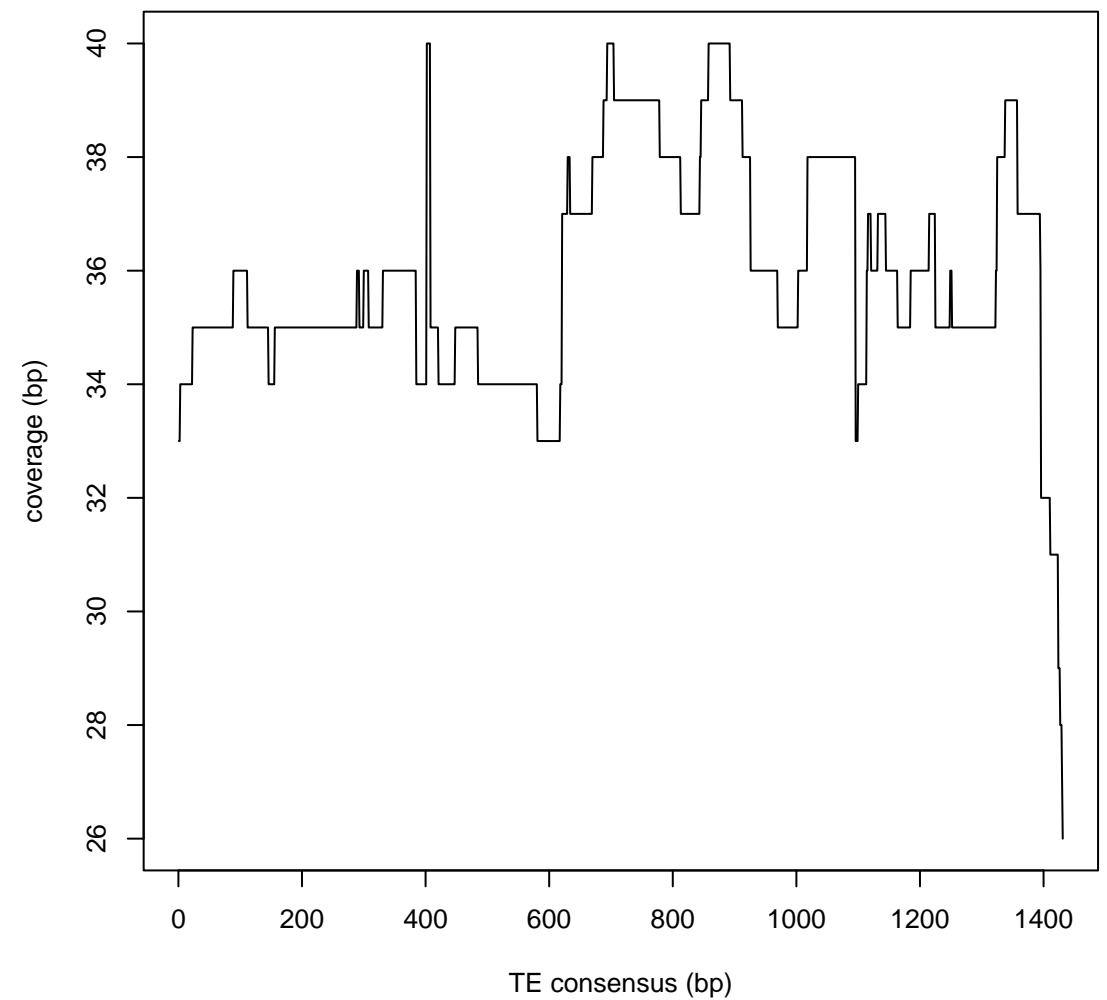


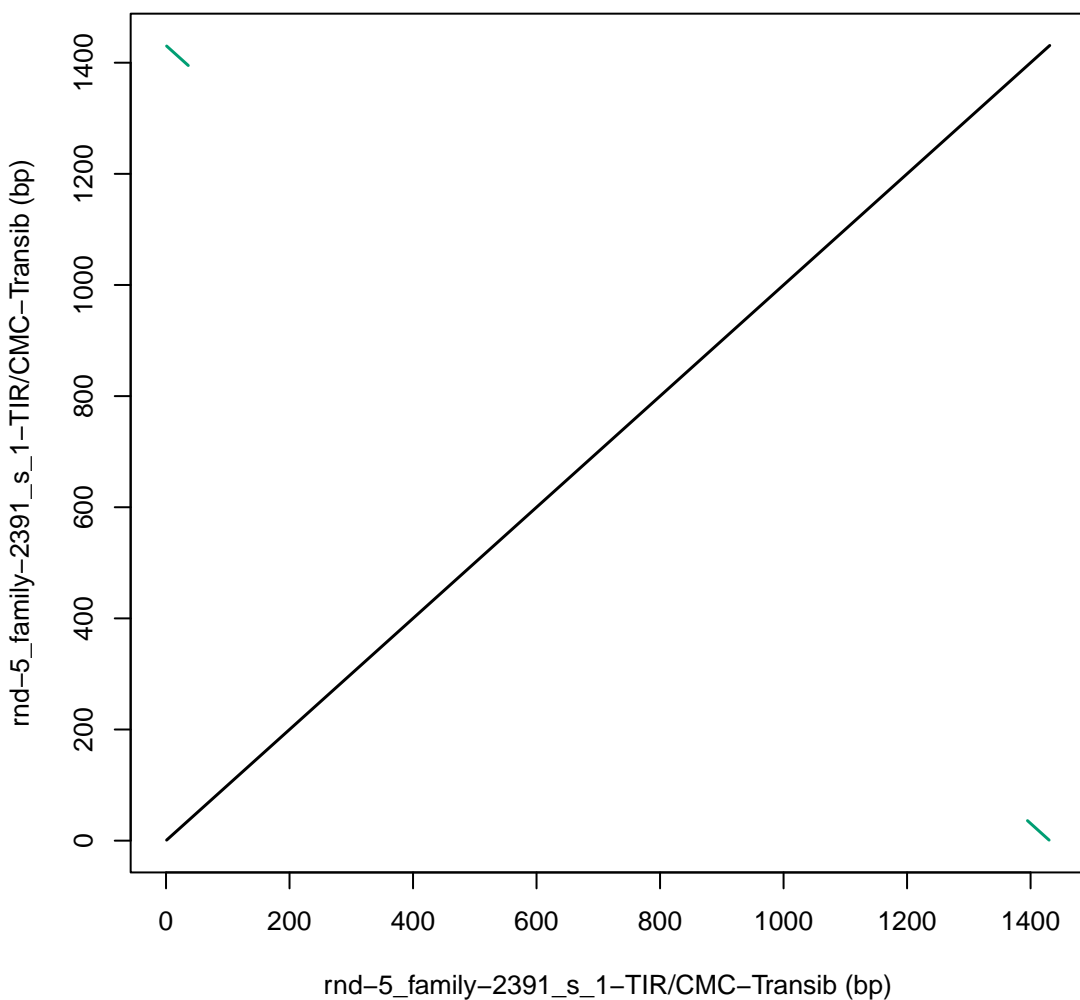
TE: rnd-5_family-2391_s_1-TIR/CMC-Transib
consensus size: 1431bp; fragments: 73; full length: 15 (≥ 1287.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

