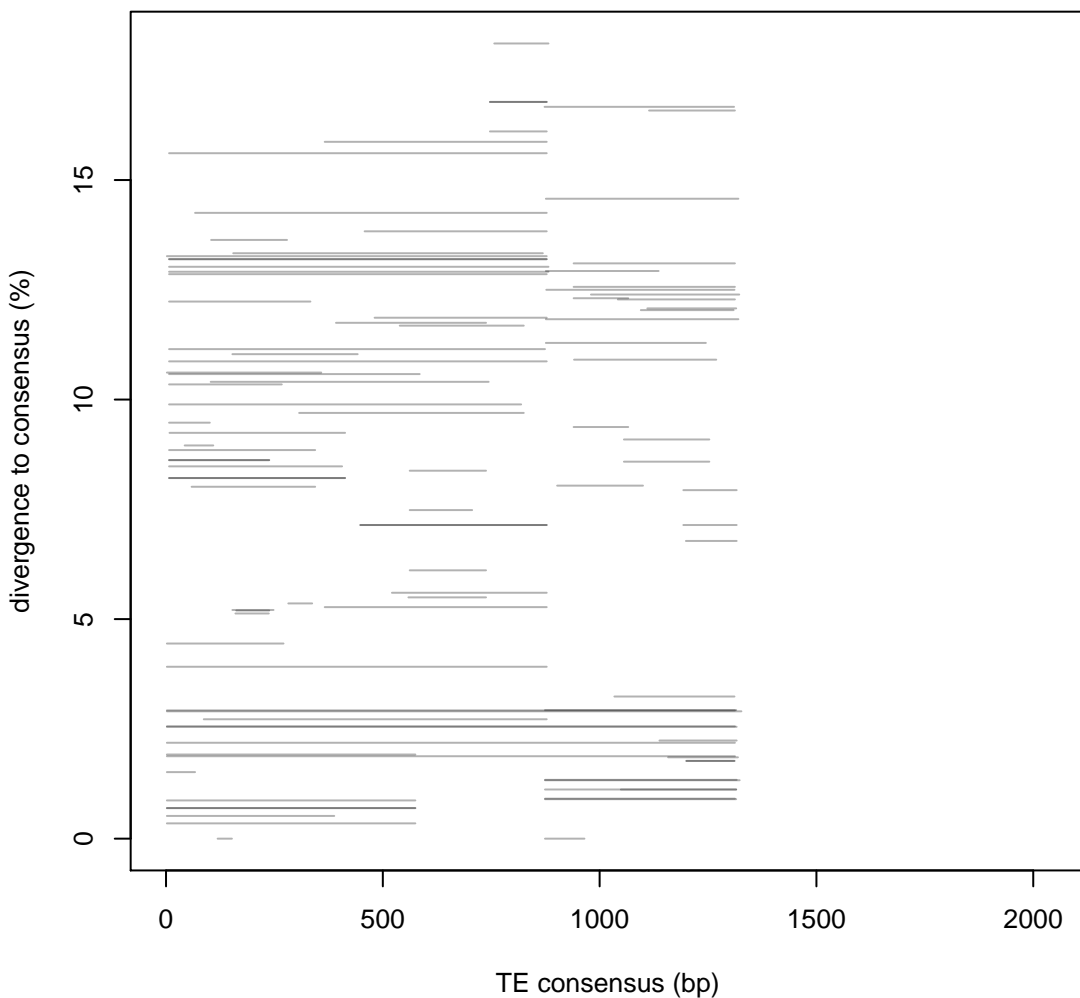
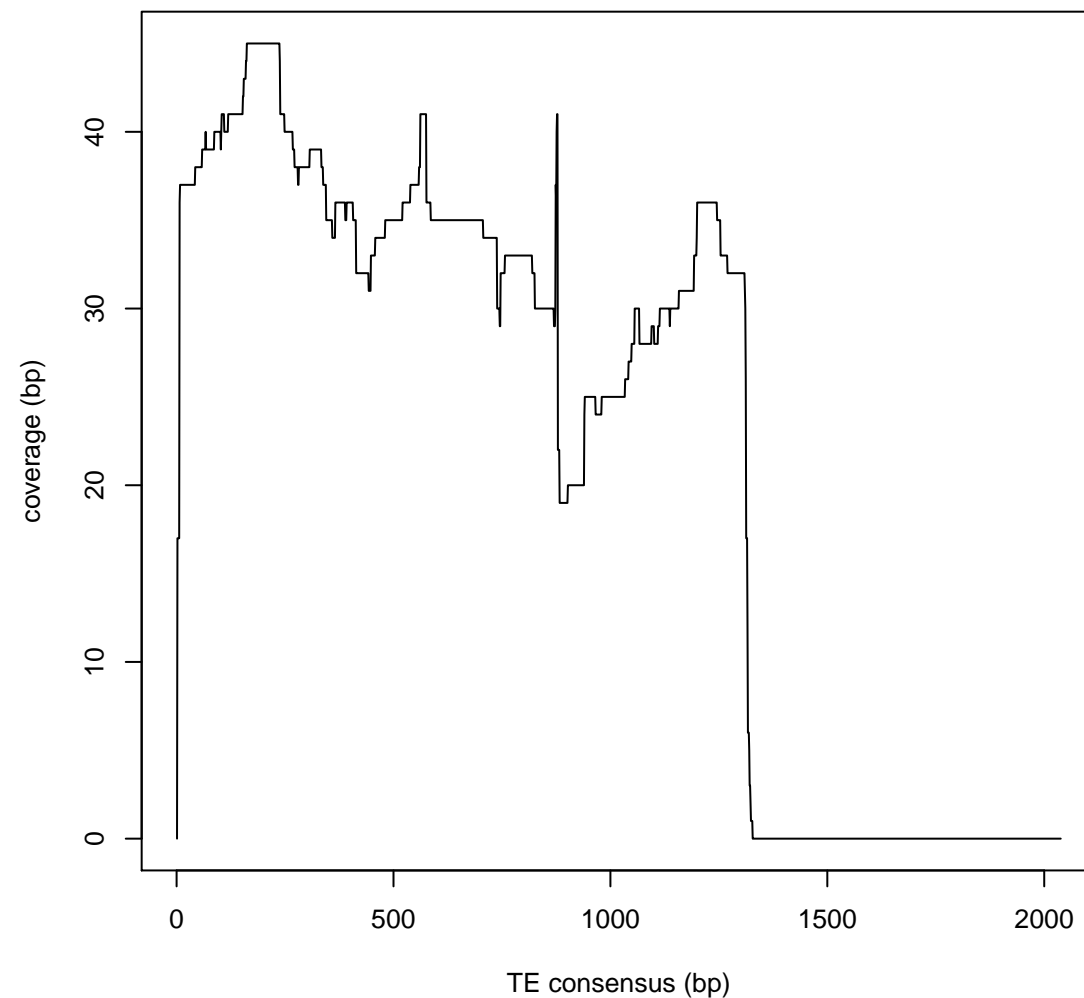


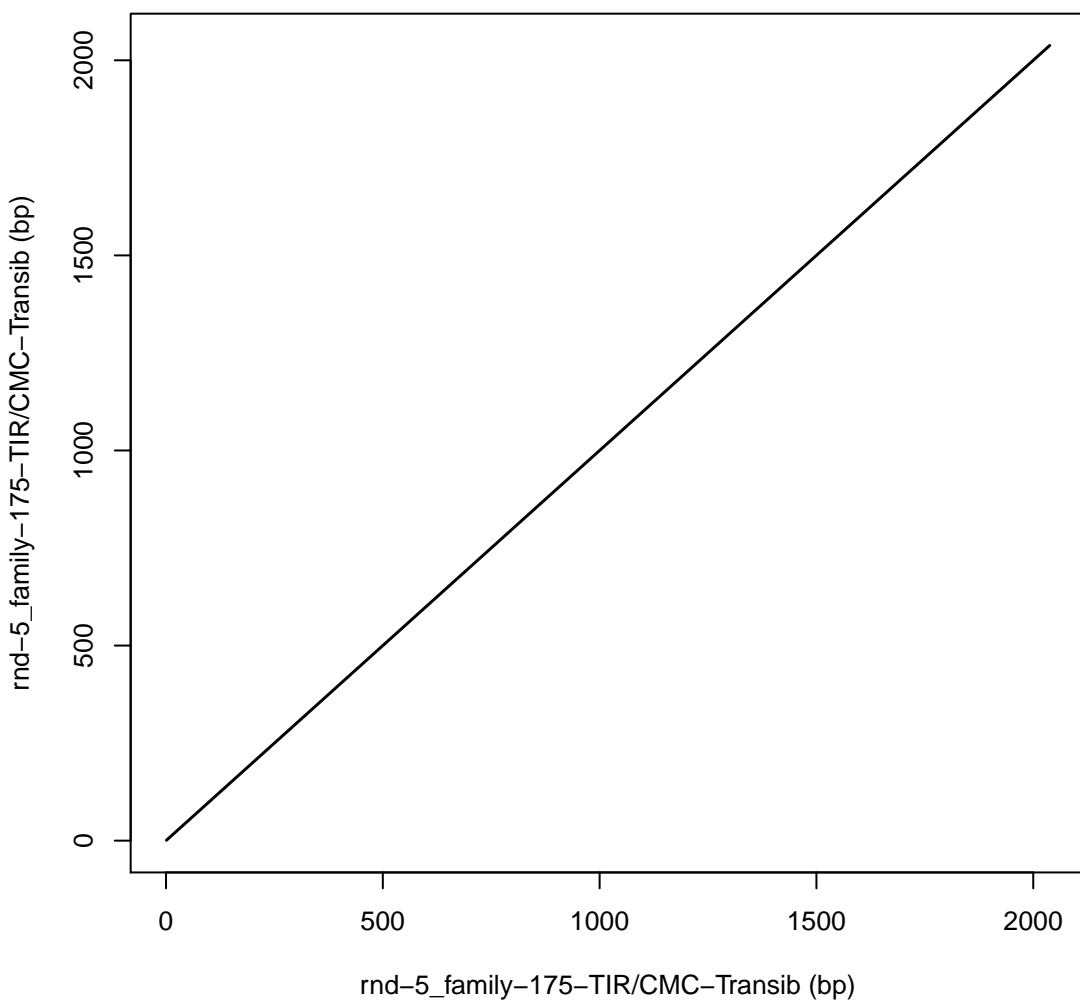
TE: rnd-5_family-175-TIR/CMC-Transib
consensus size: 2038bp; fragments: 103; full length: 0 (>=1834.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

