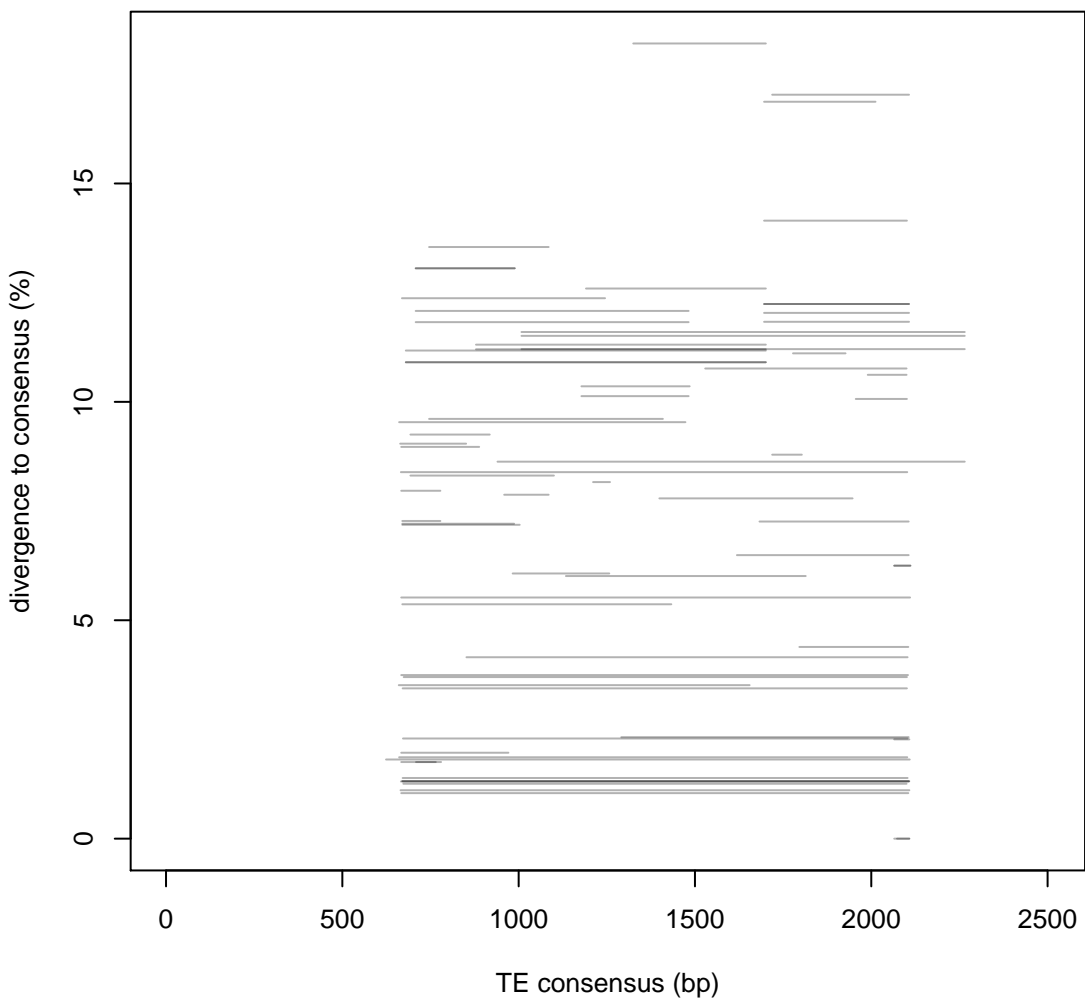
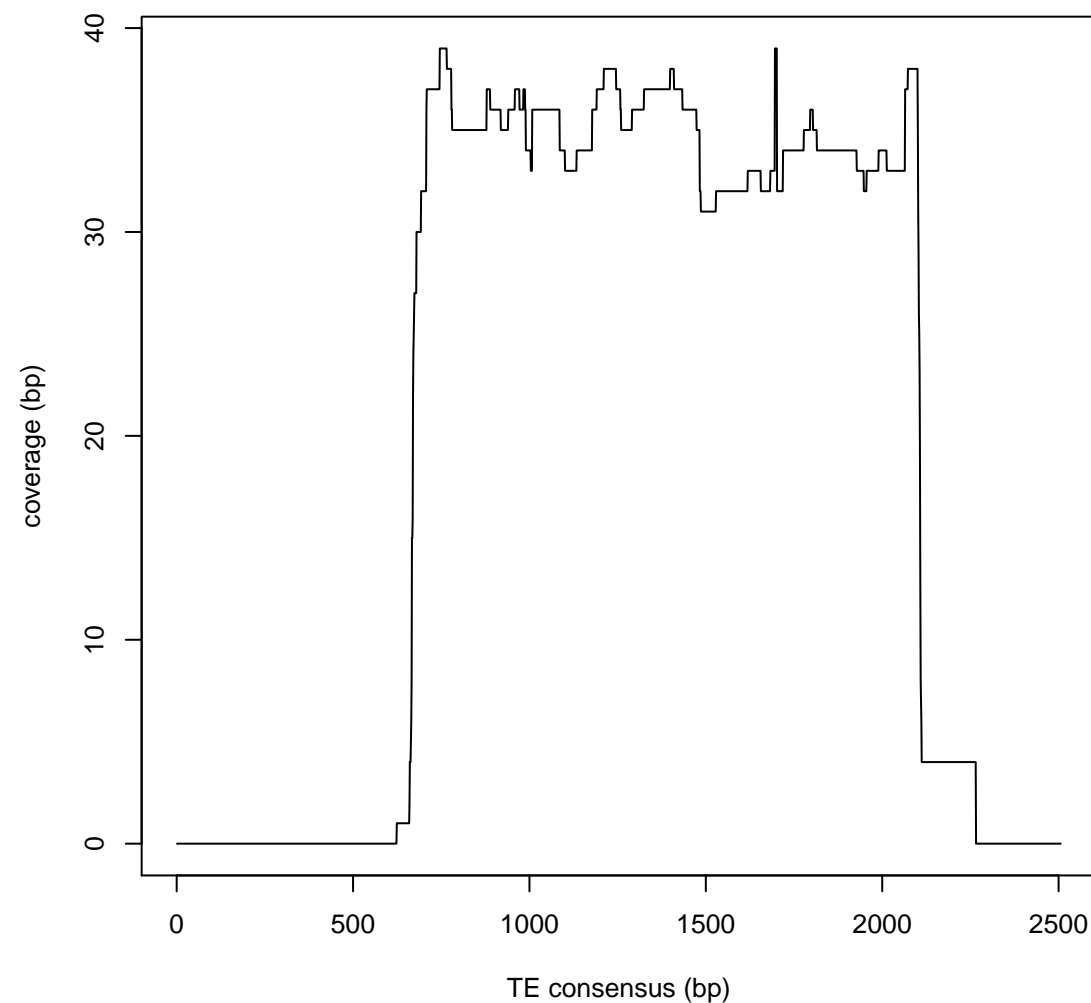


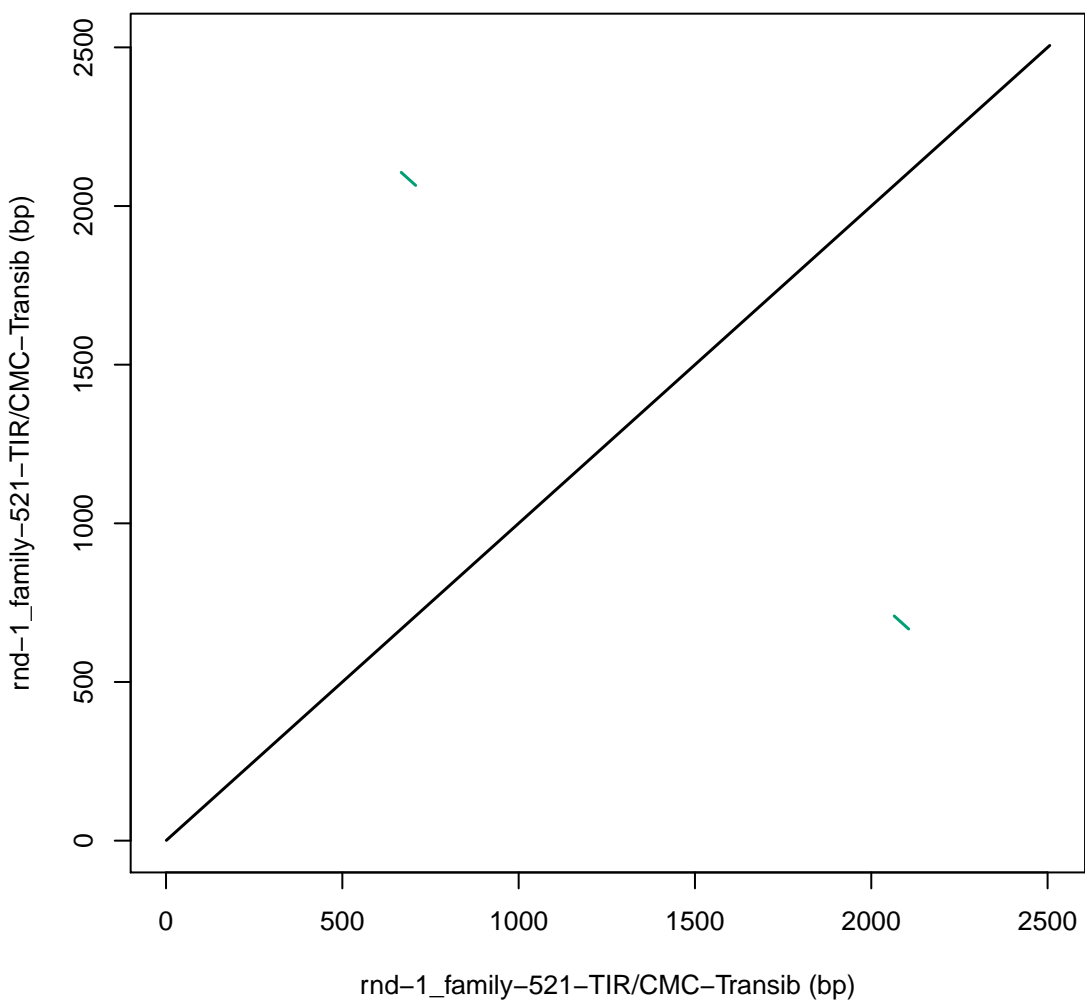
TE: rnd-1_family-521-TIR/CMC-Transib
consensus size: 2506bp; fragments: 76; full length: 0 (>=2255.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

