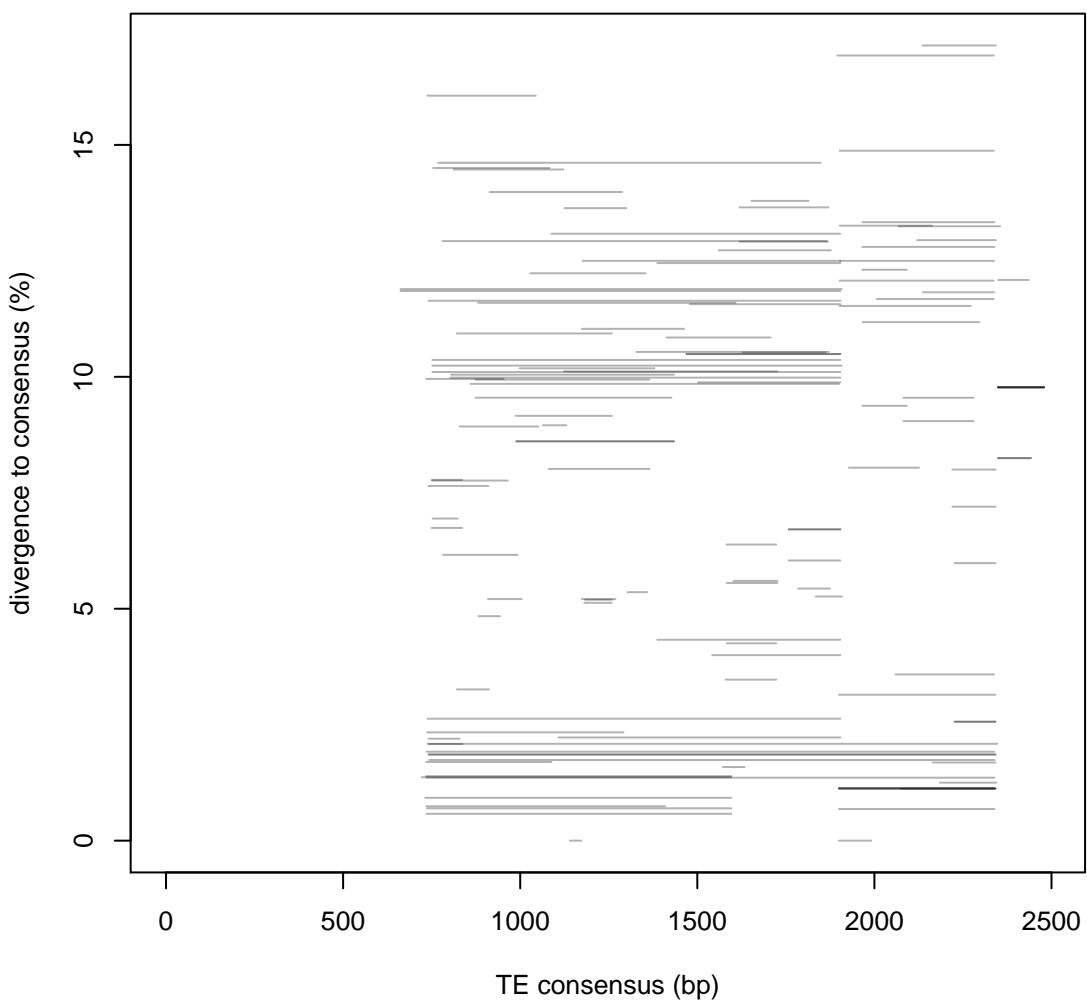
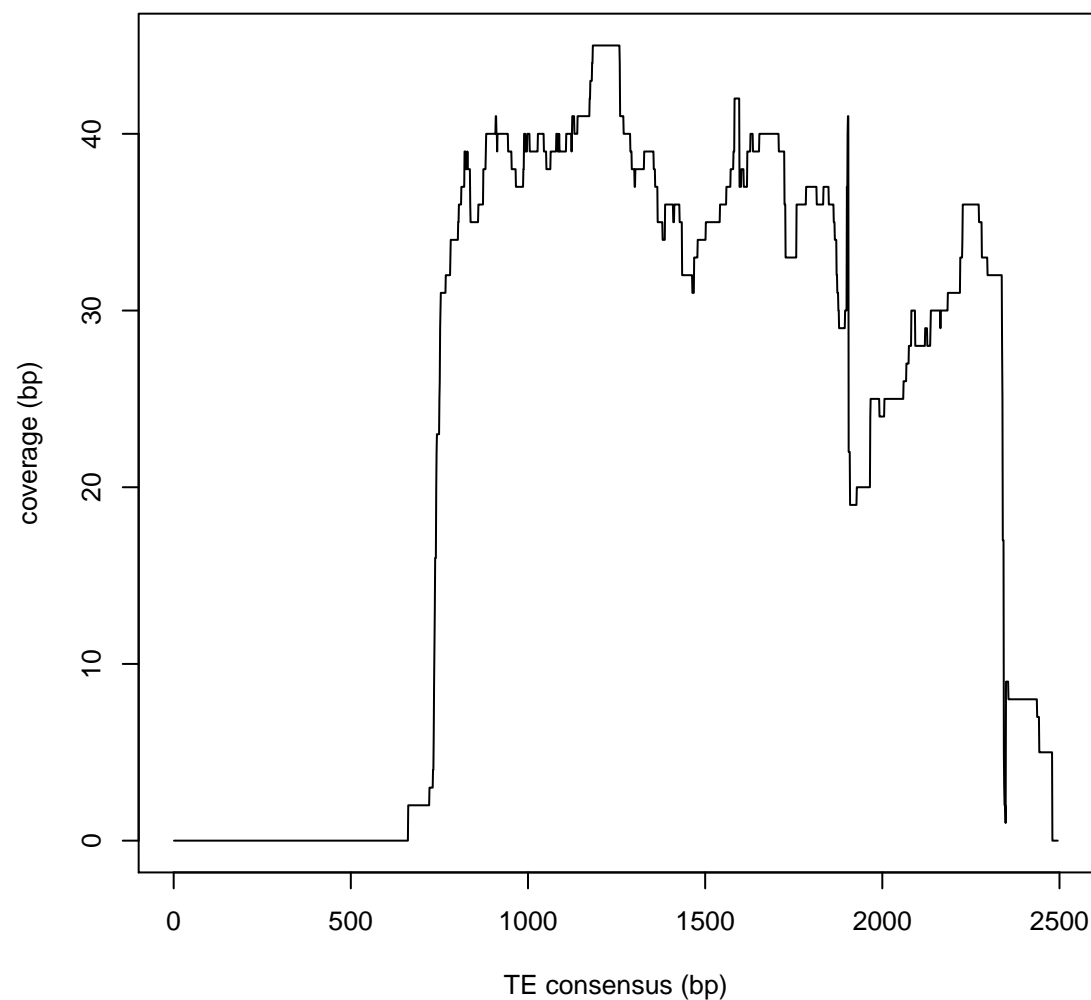


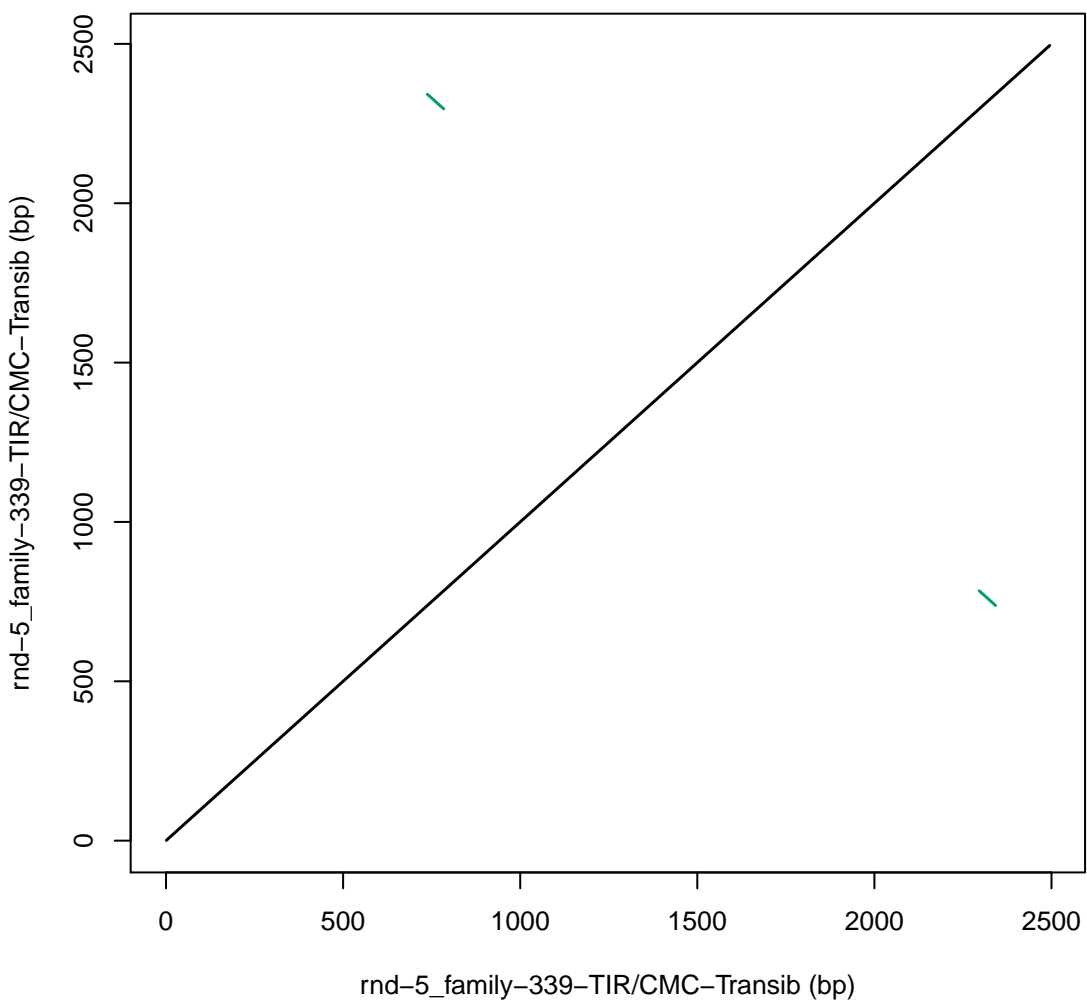
TE: rnd-5_family-339-TIR/CMC-Transib
consensus size: 2495bp; fragments: 133; full length: 0 (>=2245.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

