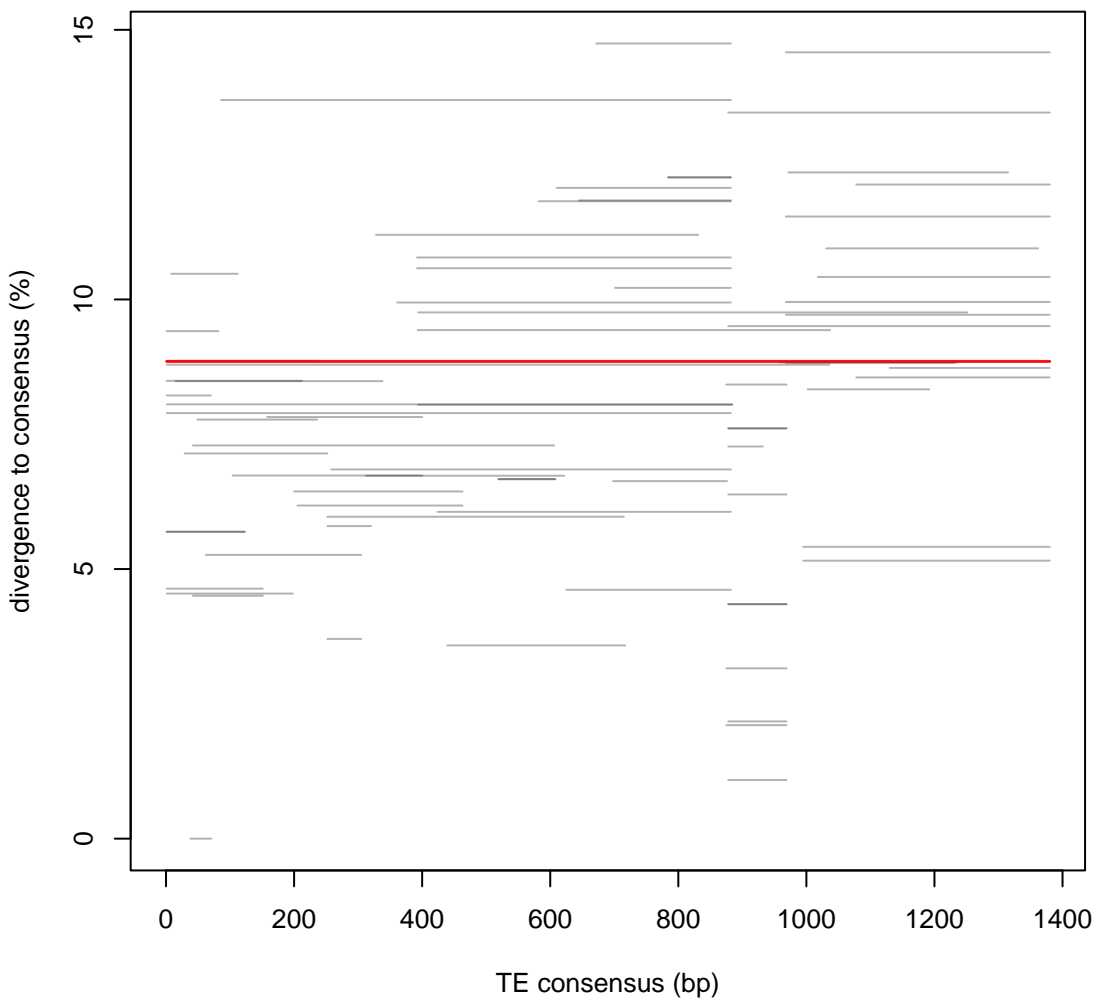
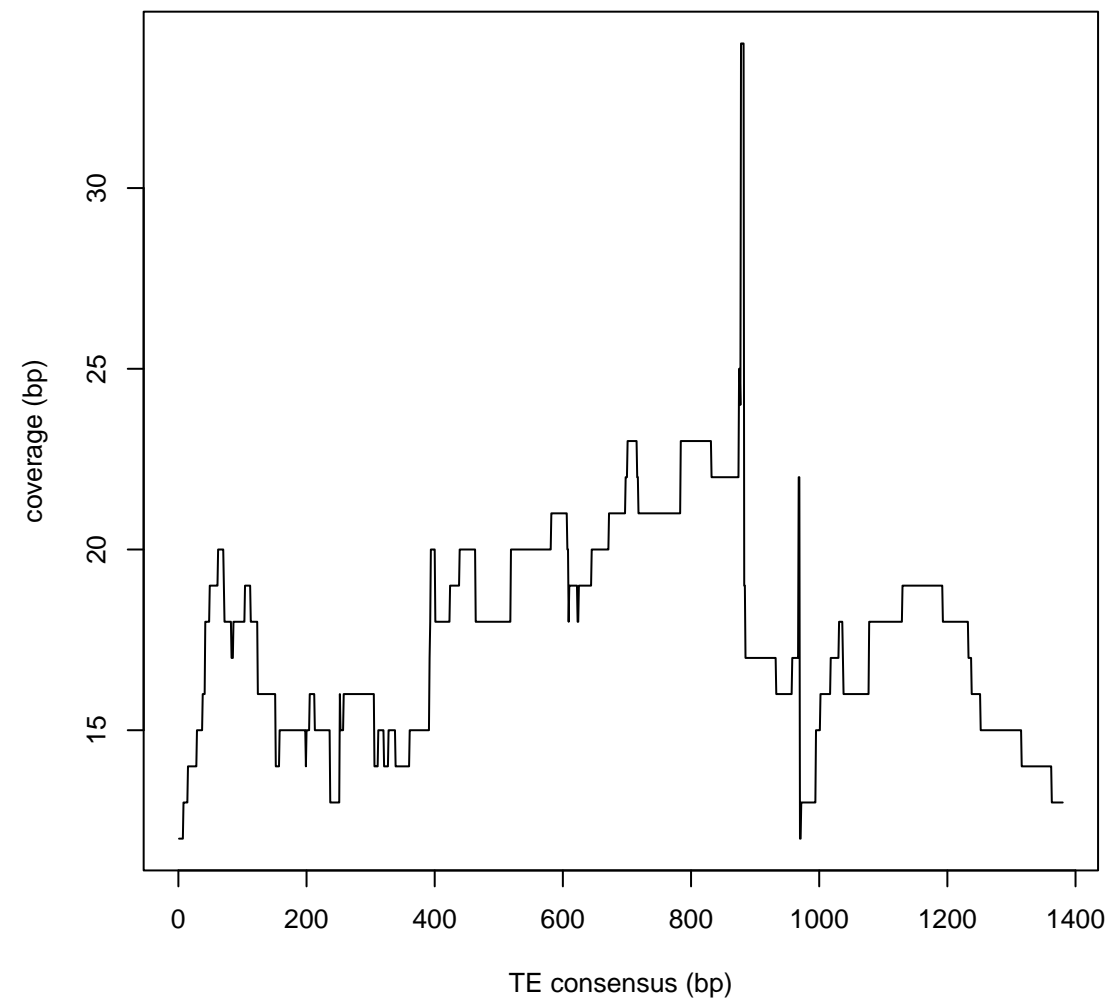


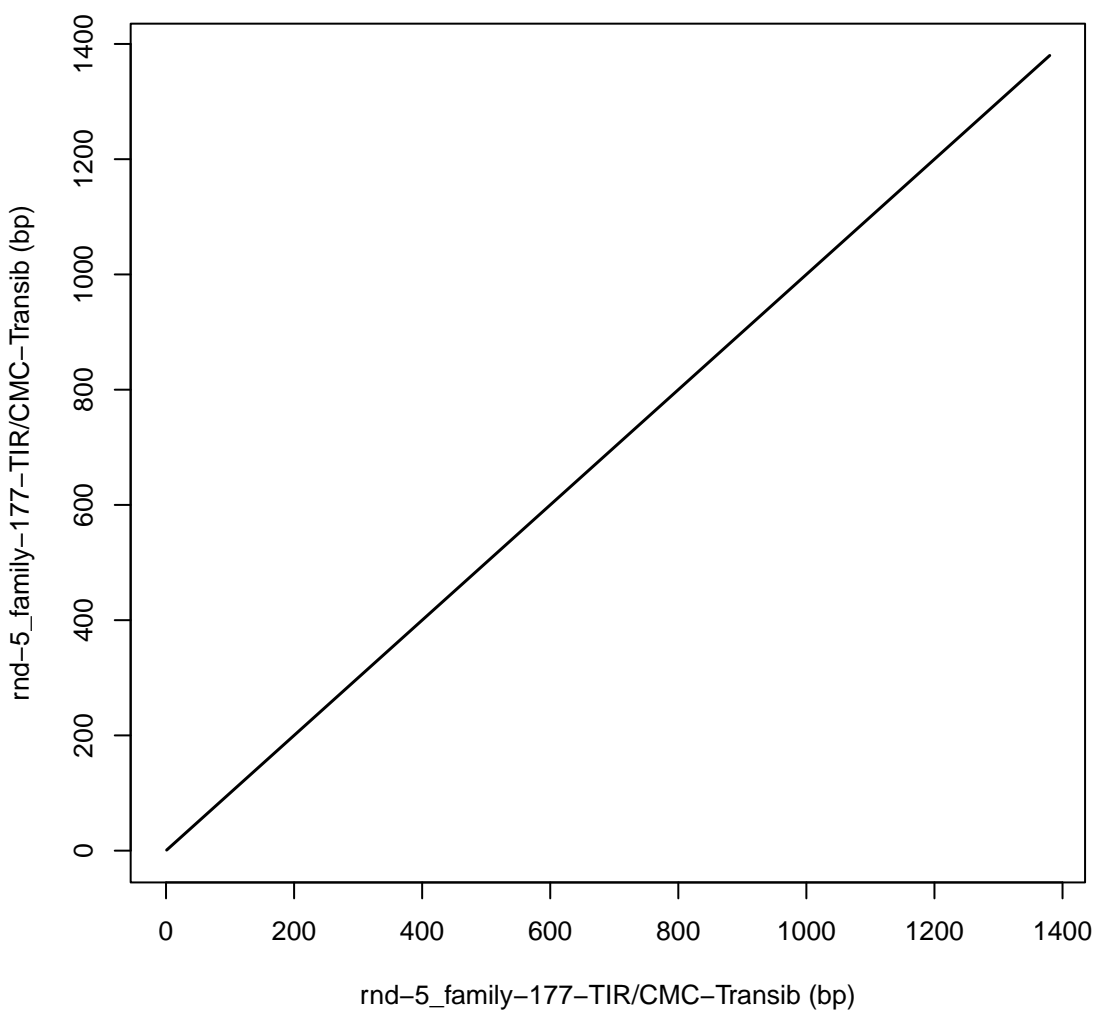
TE: rnd-5_family-177-TIR/CMC-Transib
consensus size: 1380bp; fragments: 78; full length: 1 (>=1242bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

