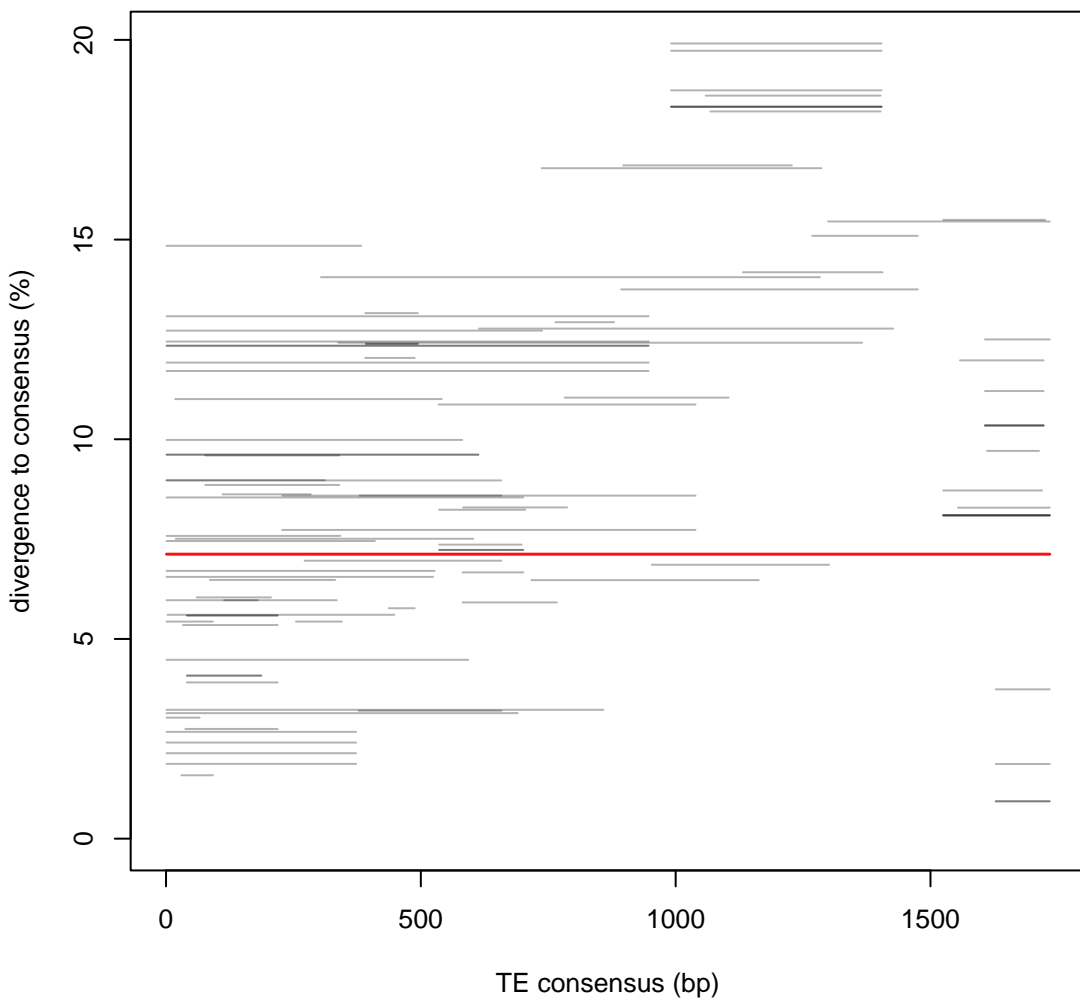
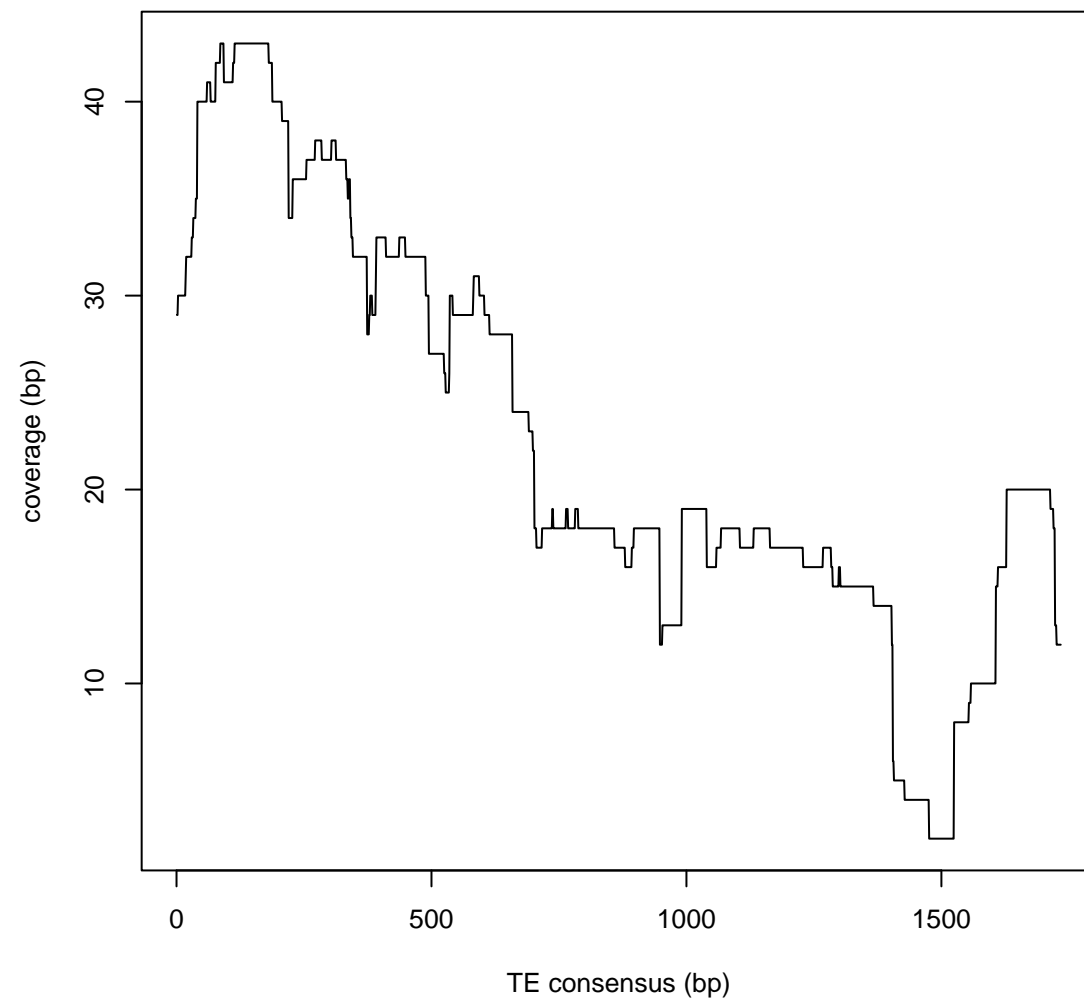


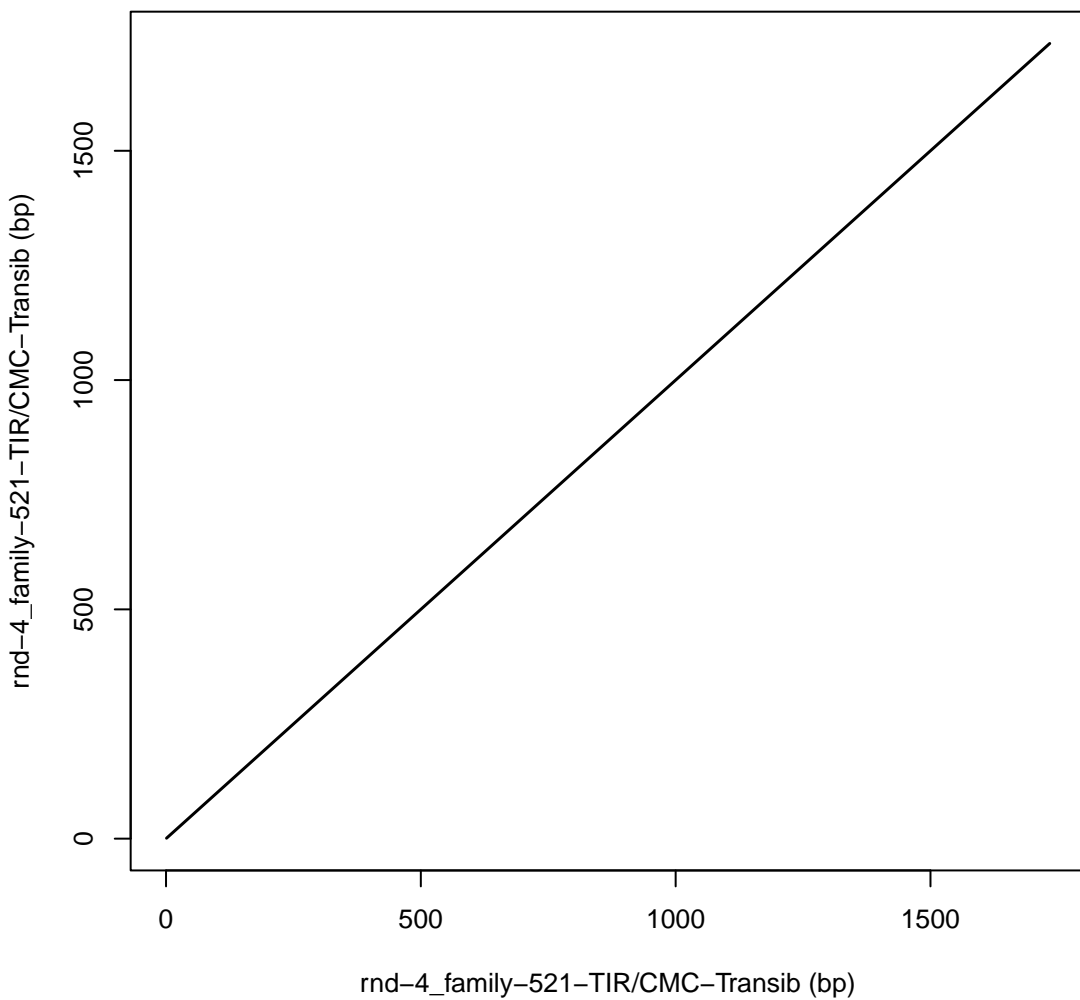
TE: rnd-4\_family-521-TIR/CMC-Transib  
consensus size: 1734bp; fragments: 104; full length: 1 (>=1560.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

