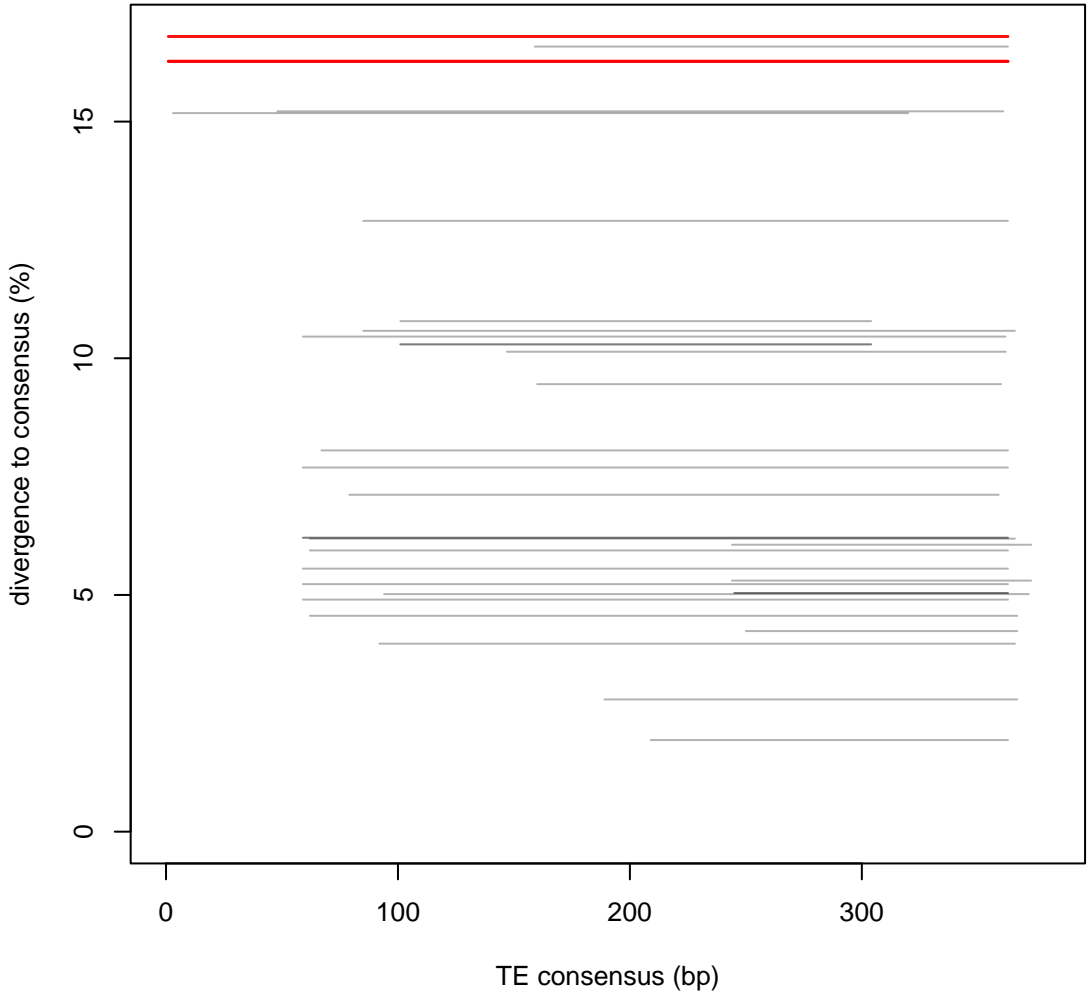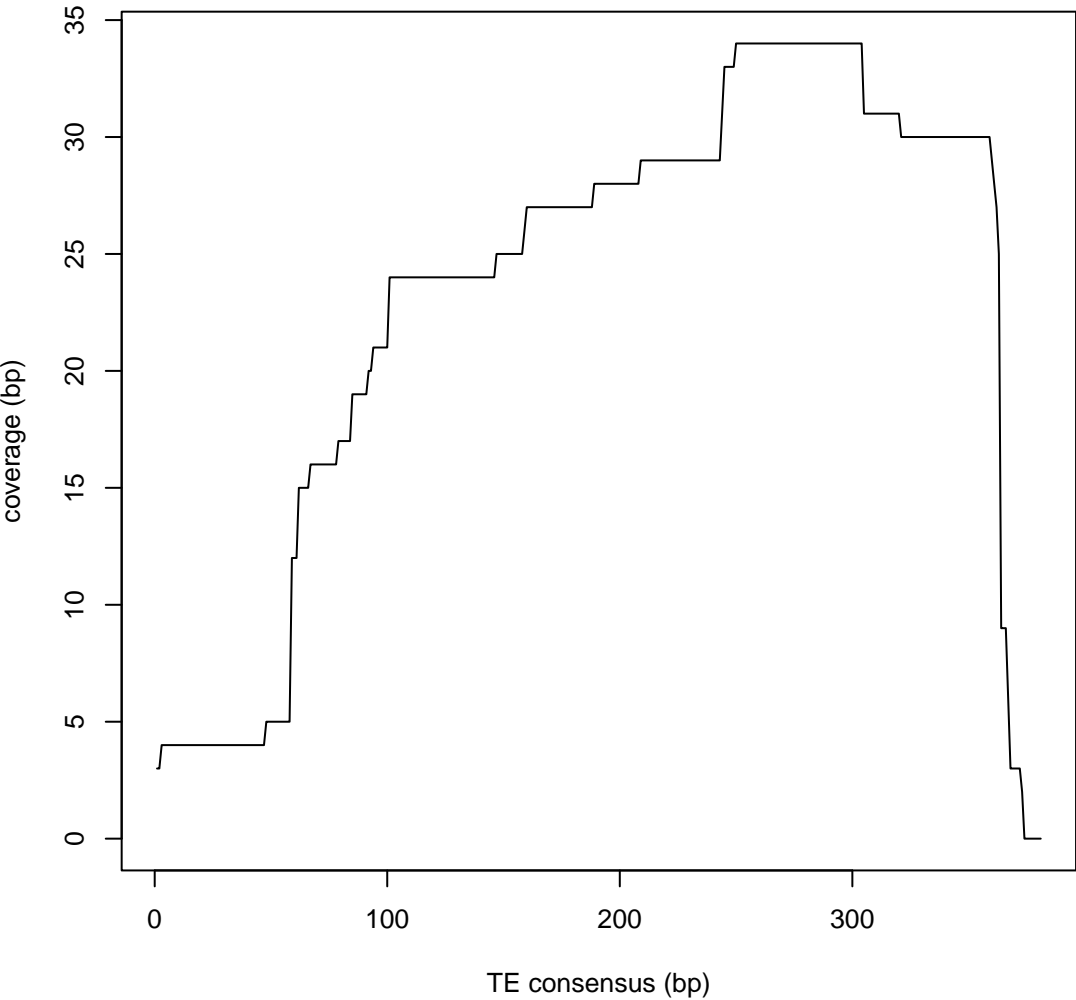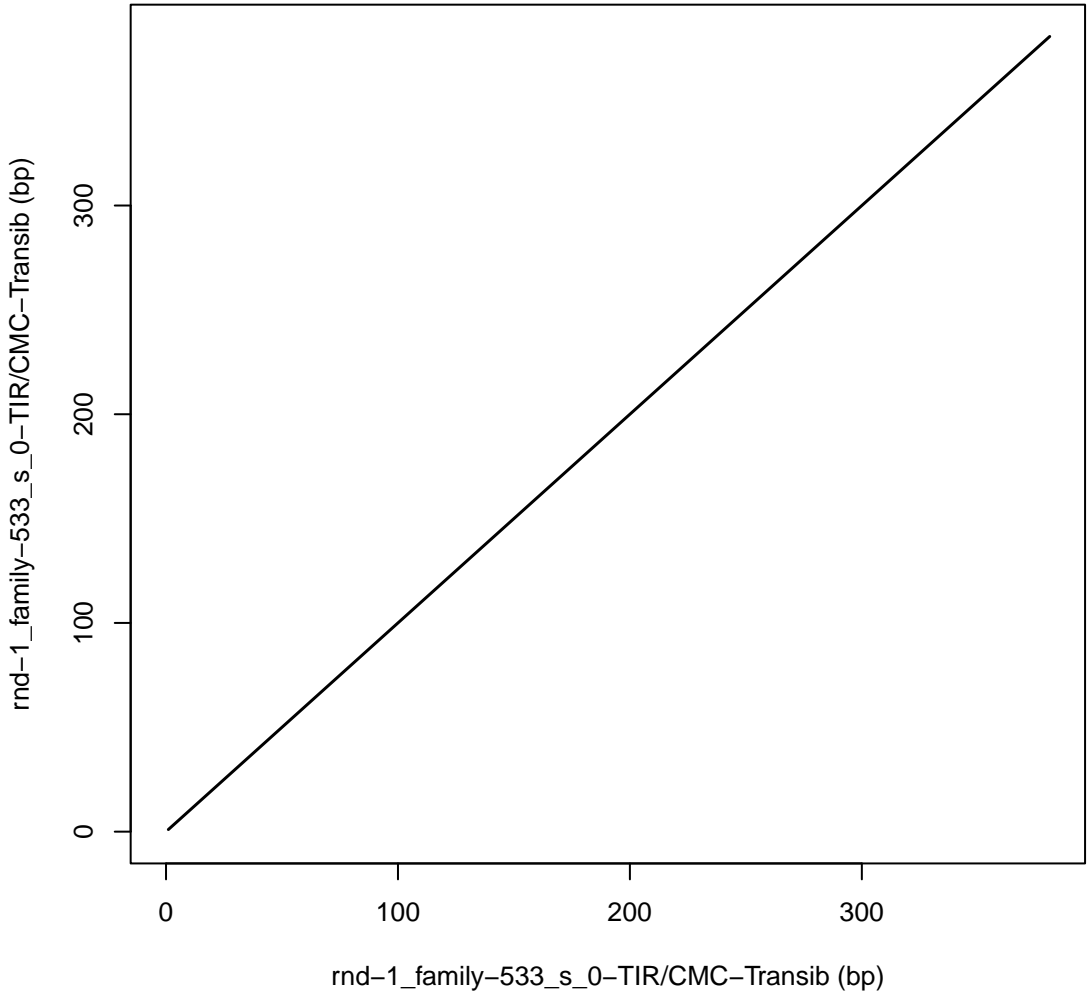**TE: rnd-1_family-533_s_0-TIR/CMC-Transib**
consensus size: 381bp; fragments: 34; full length: 3 (>=342.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected