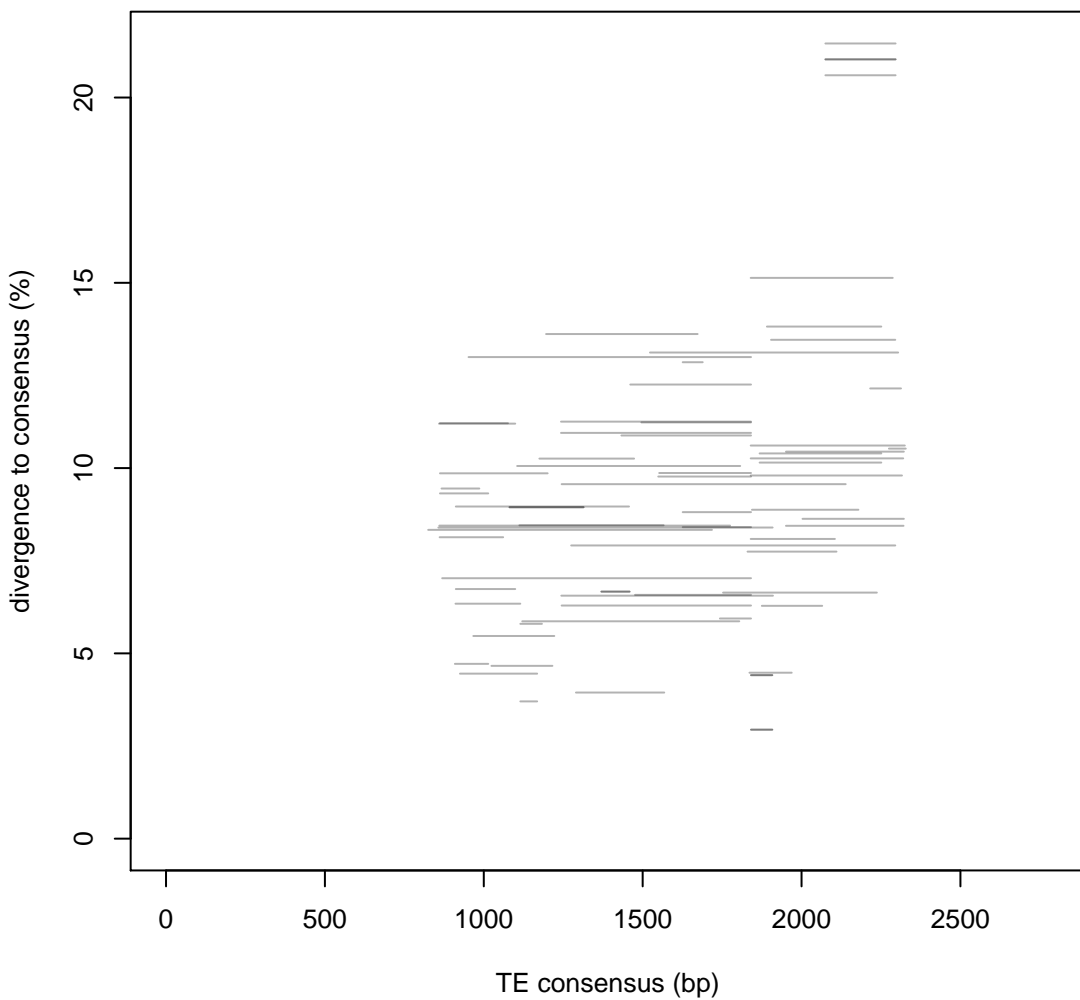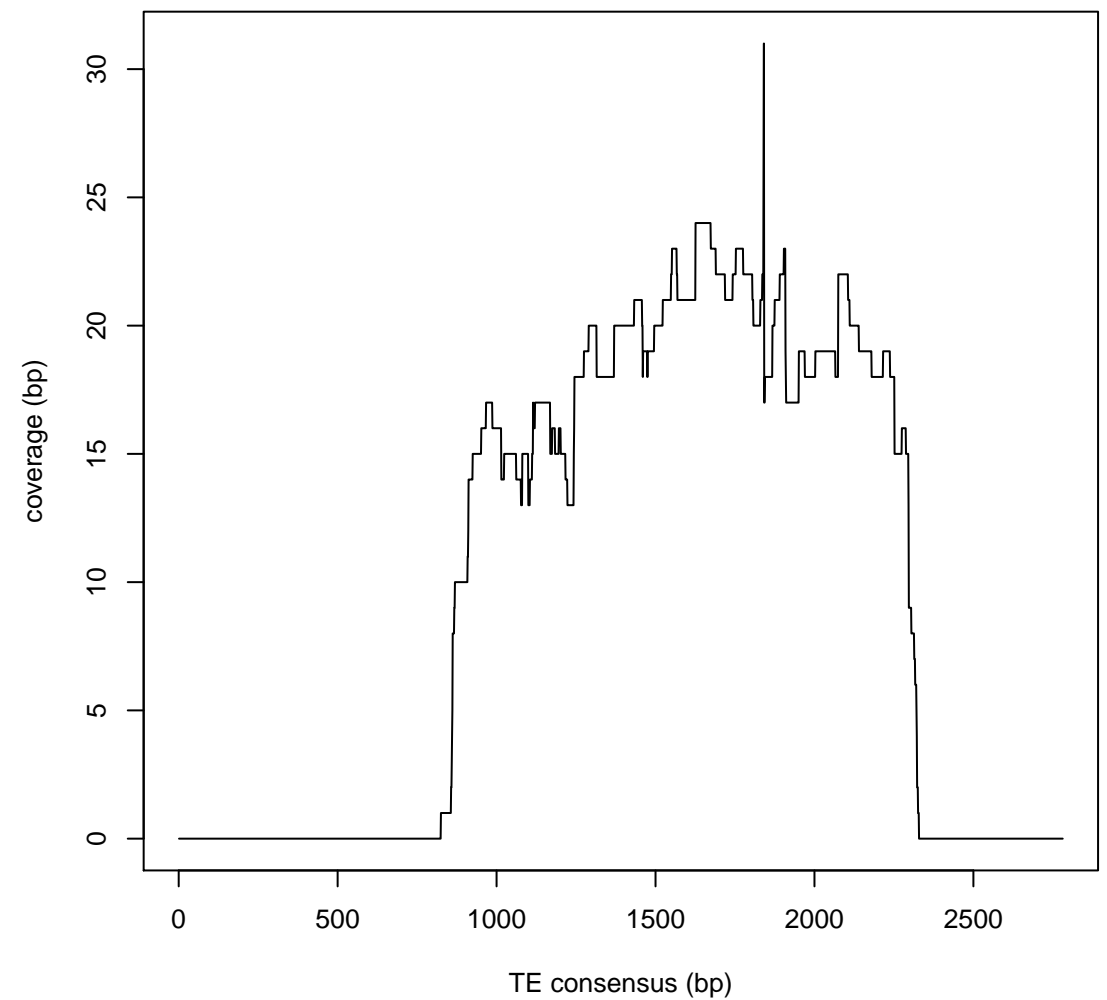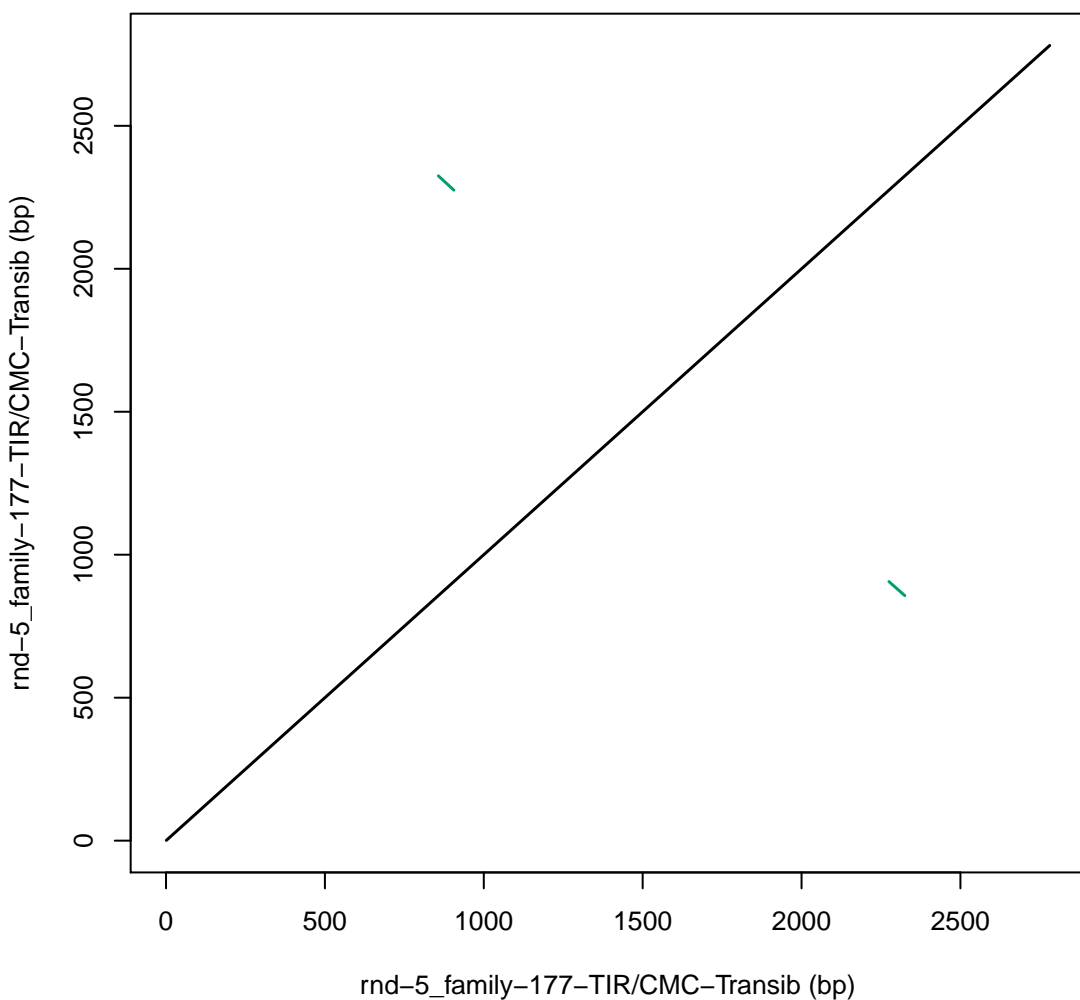**TE: rnd-5_family-177-TIR/CMC-Transib**
consensus size: 2781bp; fragments: 74; full length: 0 (>=2502.9bp)

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd-5_family-177-TIR/CMC-Transib (bp)

rnd-5_family-177-TIR/CMC-Transib (bp)

TE consensus structure and protein hits

no orf >400 bp detected

rnd-5_family-177-TIR/CMC-Transib (bp)