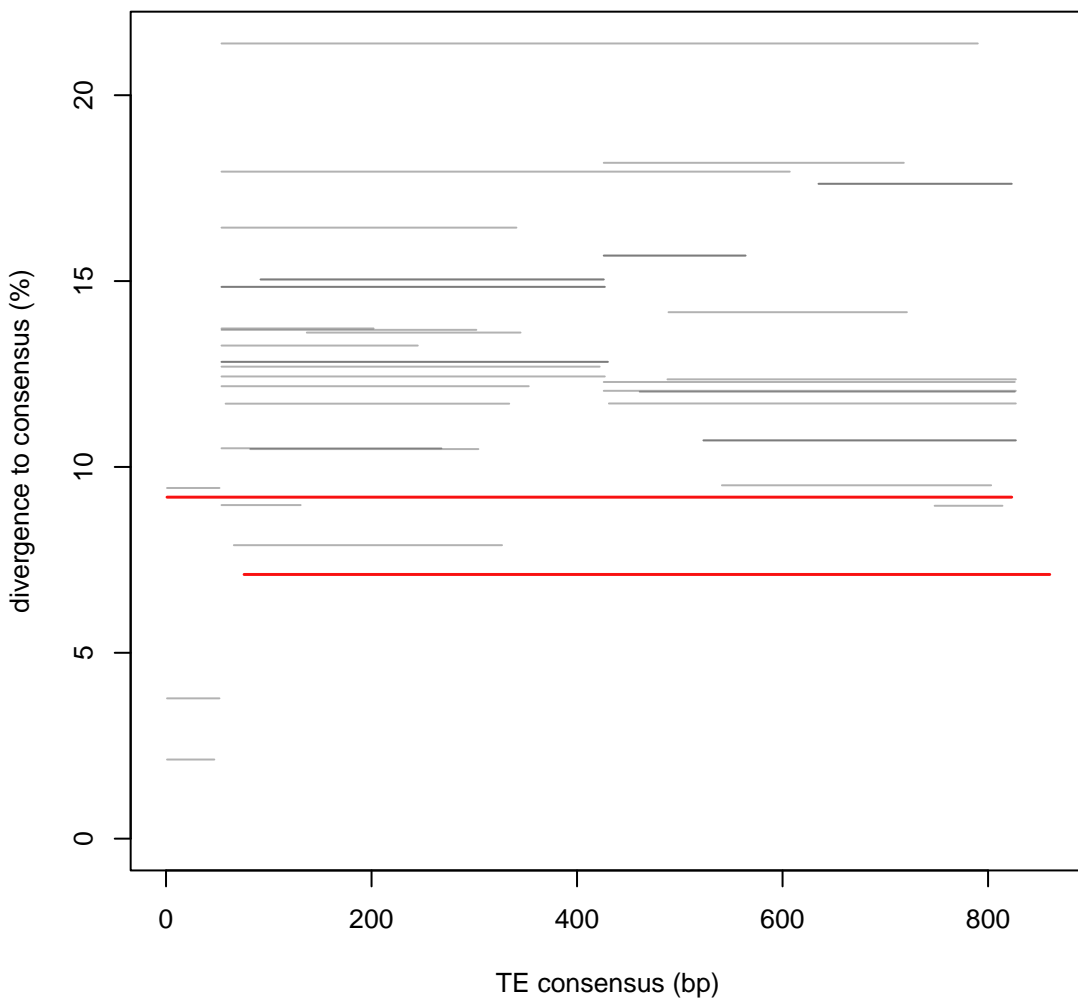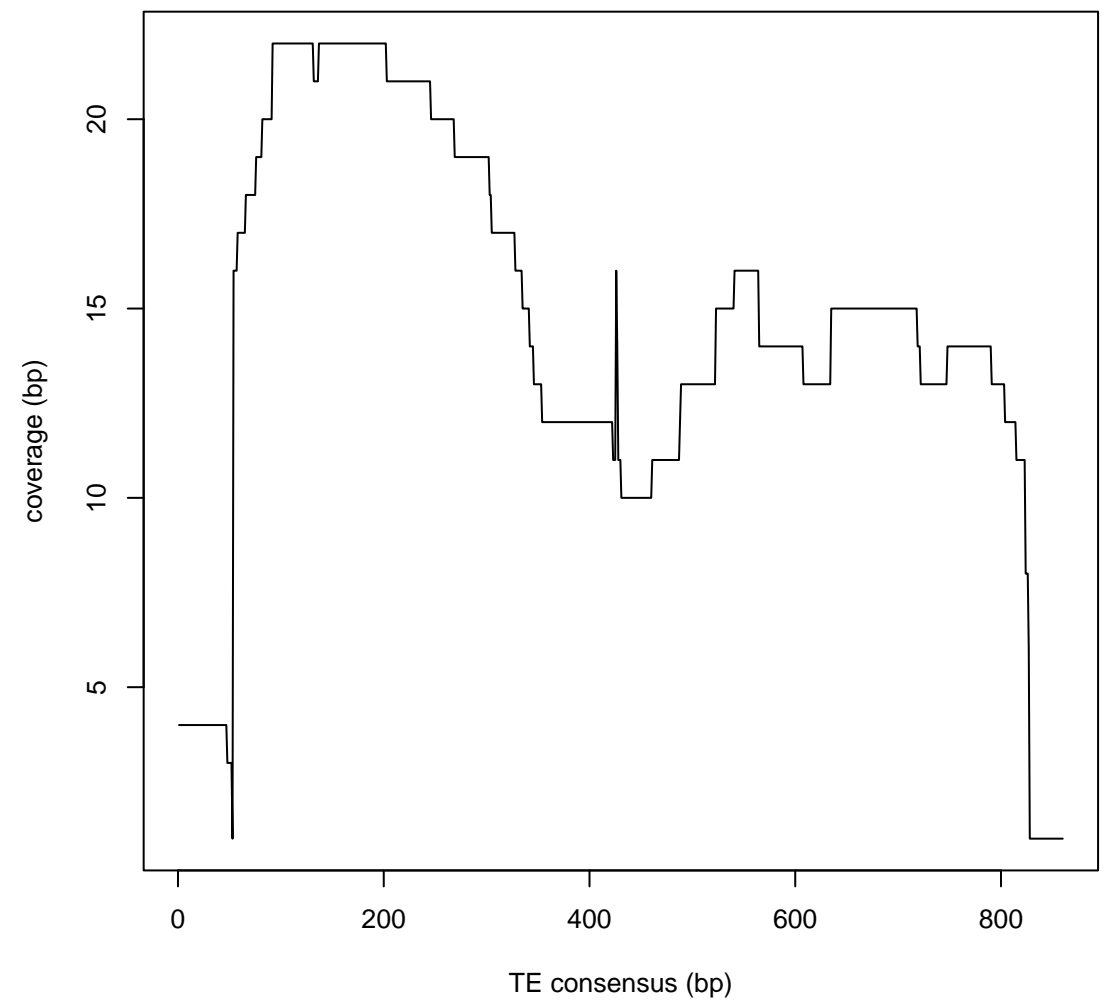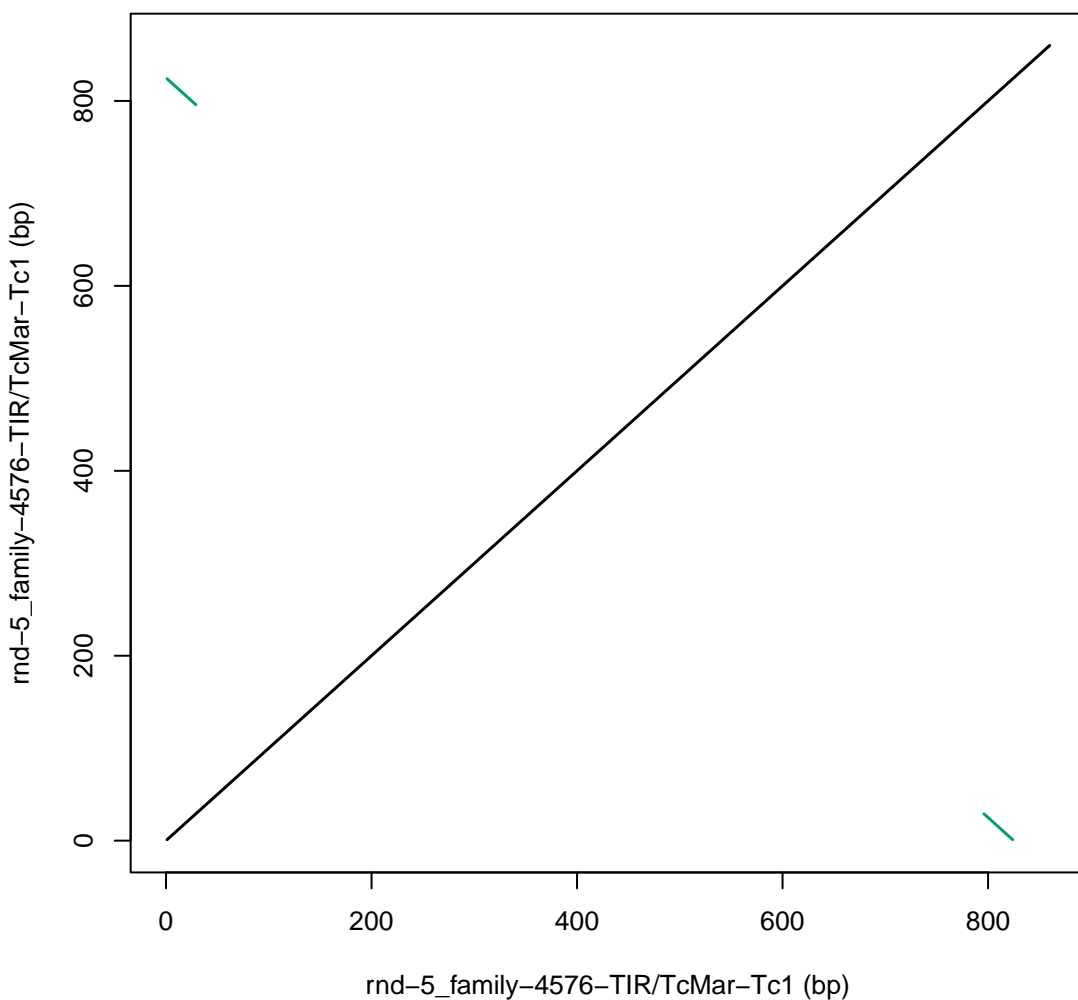## TE: rnd-5_family-4576-TIR/TcMar-Tc1
consensus size: 860bp; fragments: 41; full length: 2 (>=774bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

rnd-5_family-4576-TIR/TcMar-Tc1 (bp)

rnd-5_family-4576-TIR/TcMar-Tc1 (bp)

rnd-5_family-4576-TIR/TcMar-Tc1 (bp)