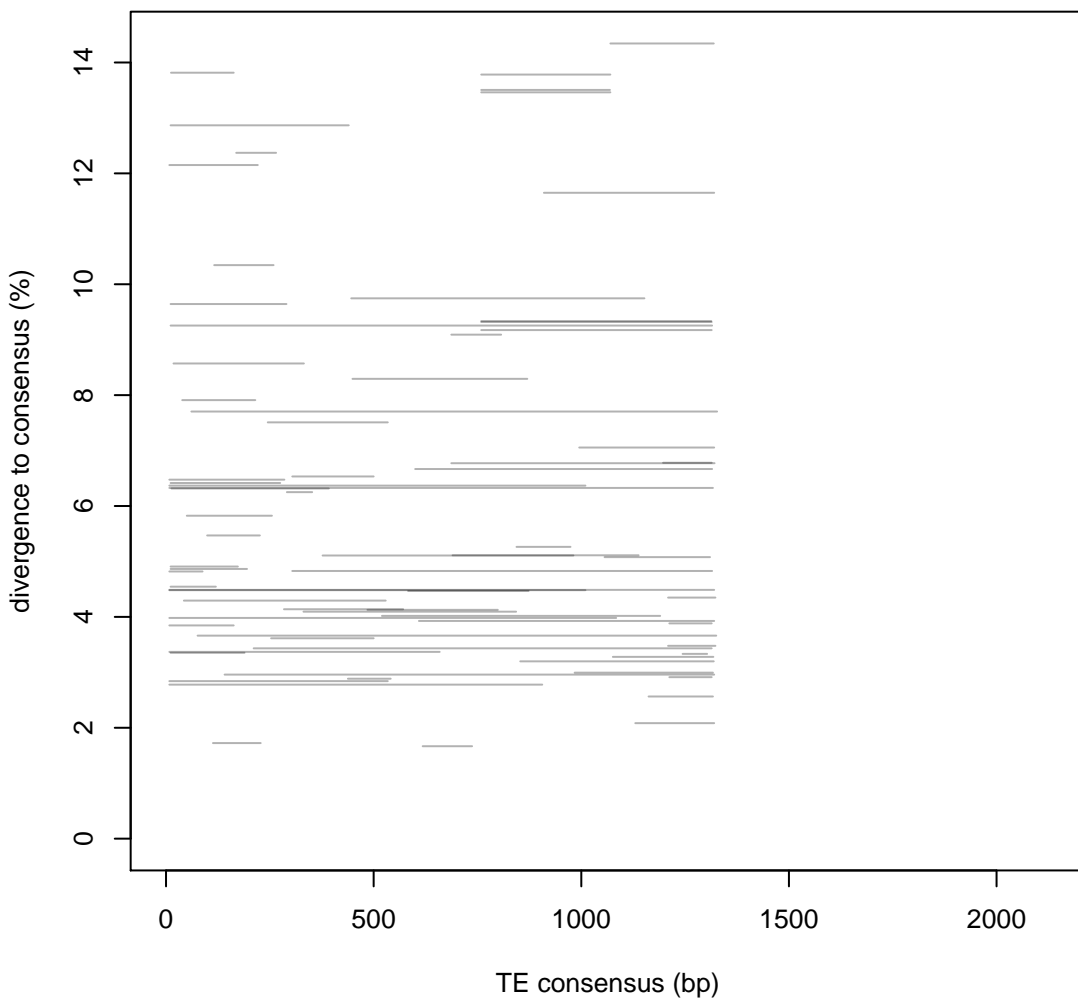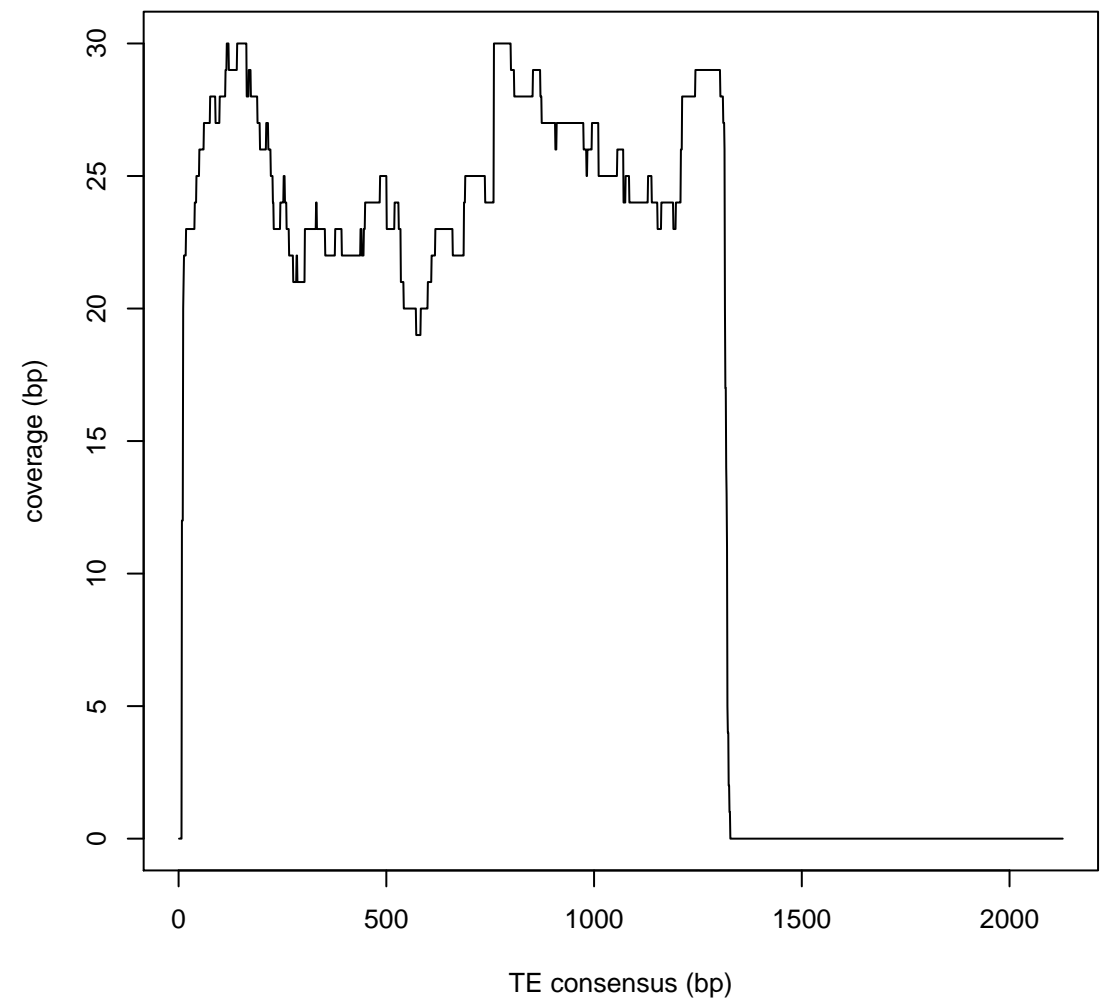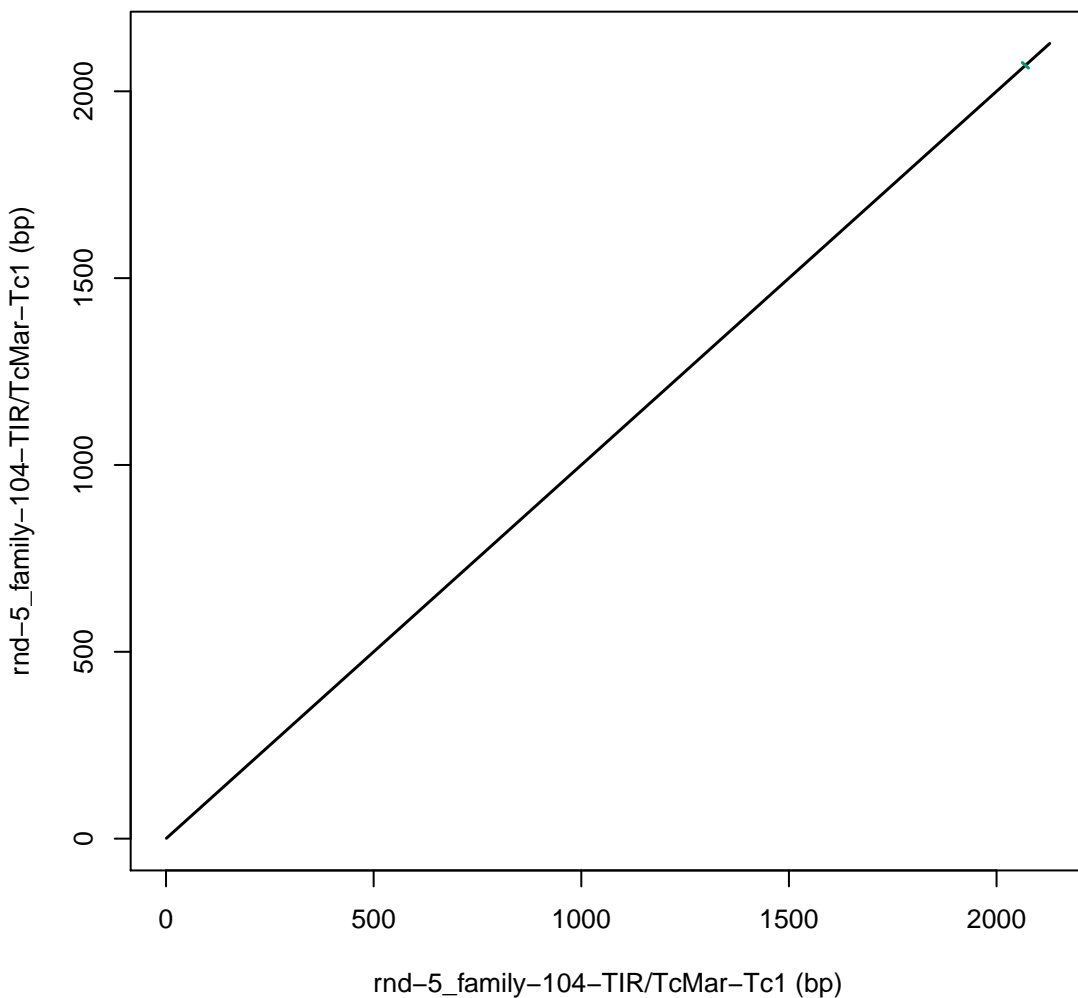**TE: rnd−5_family−104−TIR/TcMar−Tc1**
consensus size: 2128bp; fragments: 76; full length: 0 (>=1915.2bp)

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−5_family−104−TIR/TcMar−Tc1 (bp)

rnd−5_family−104−TIR/TcMar−Tc1 (bp)

TE consensus structure and protein hits

no orf >400 bp detected

rnd−5_family−104−TIR/TcMar−Tc1 (bp)