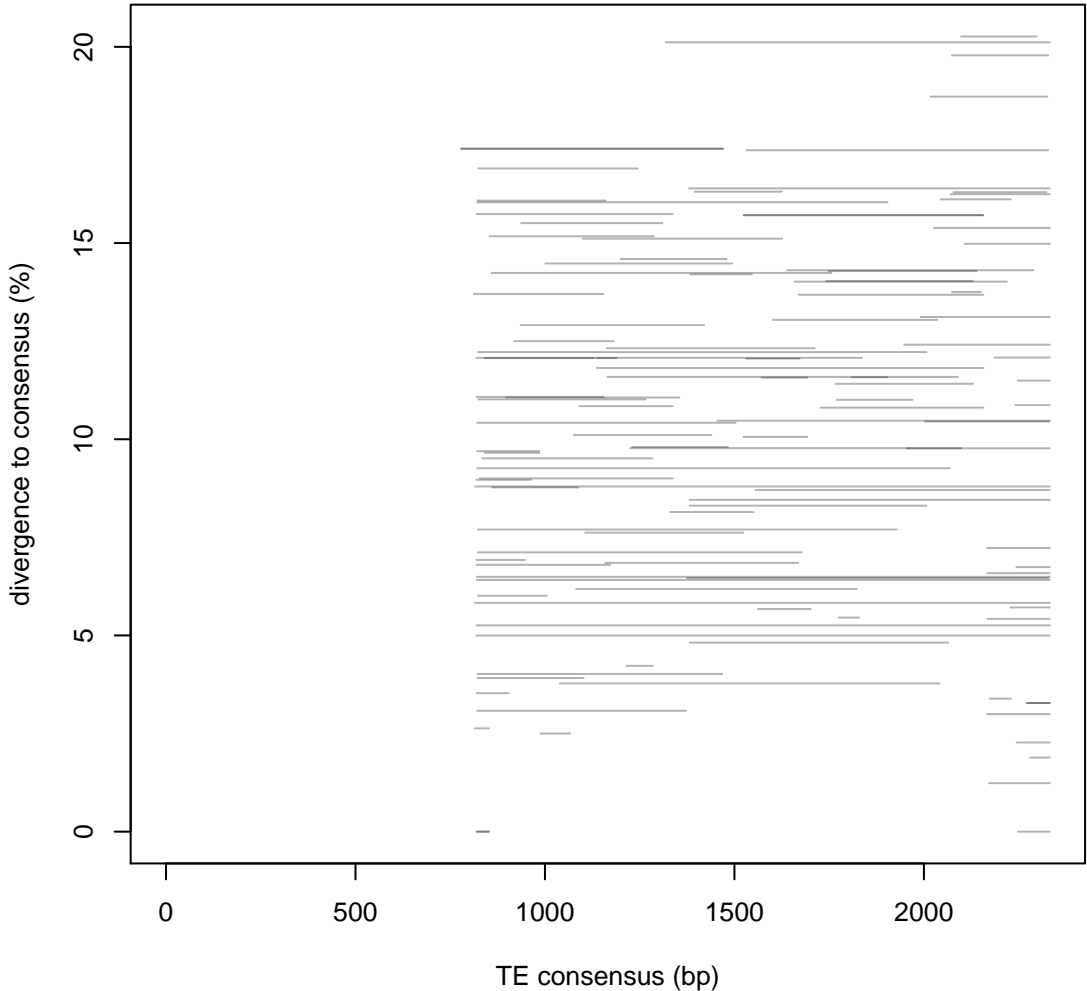
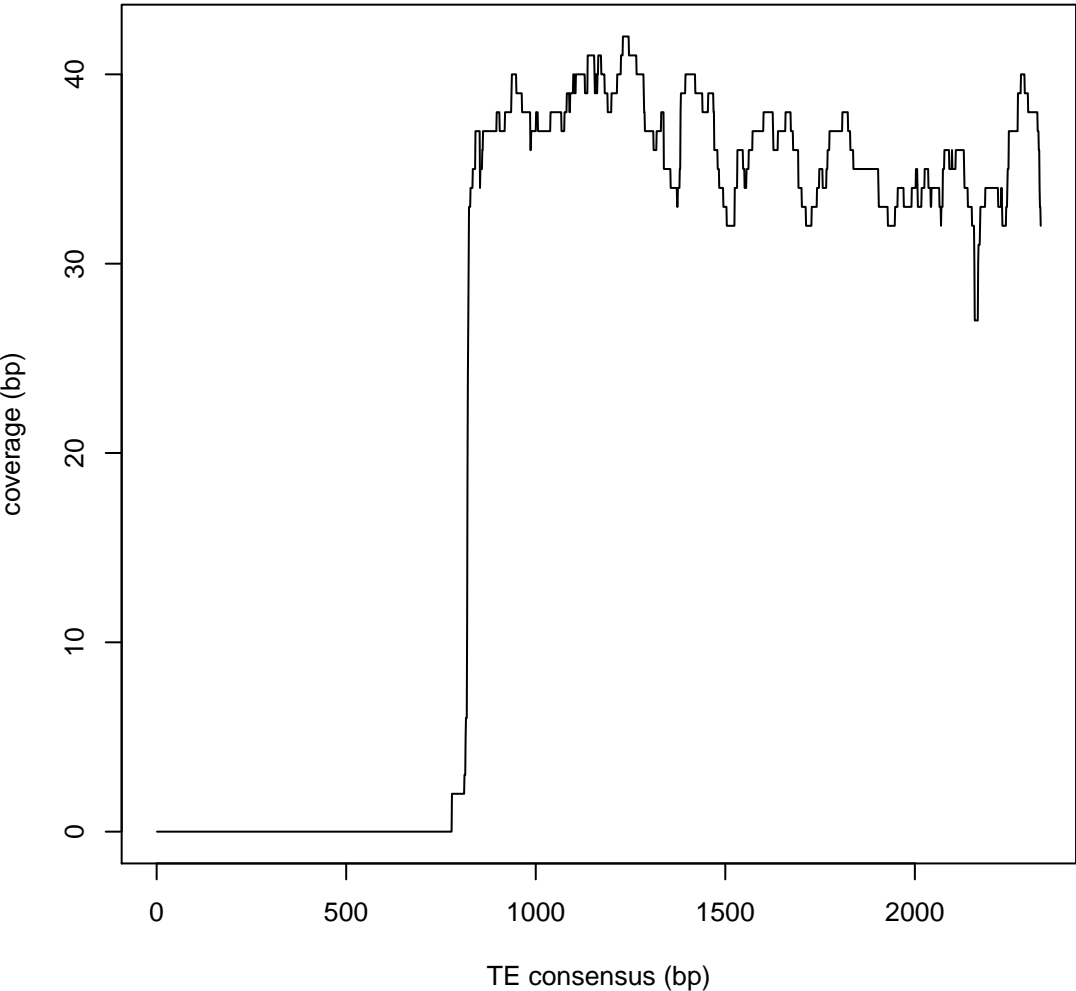


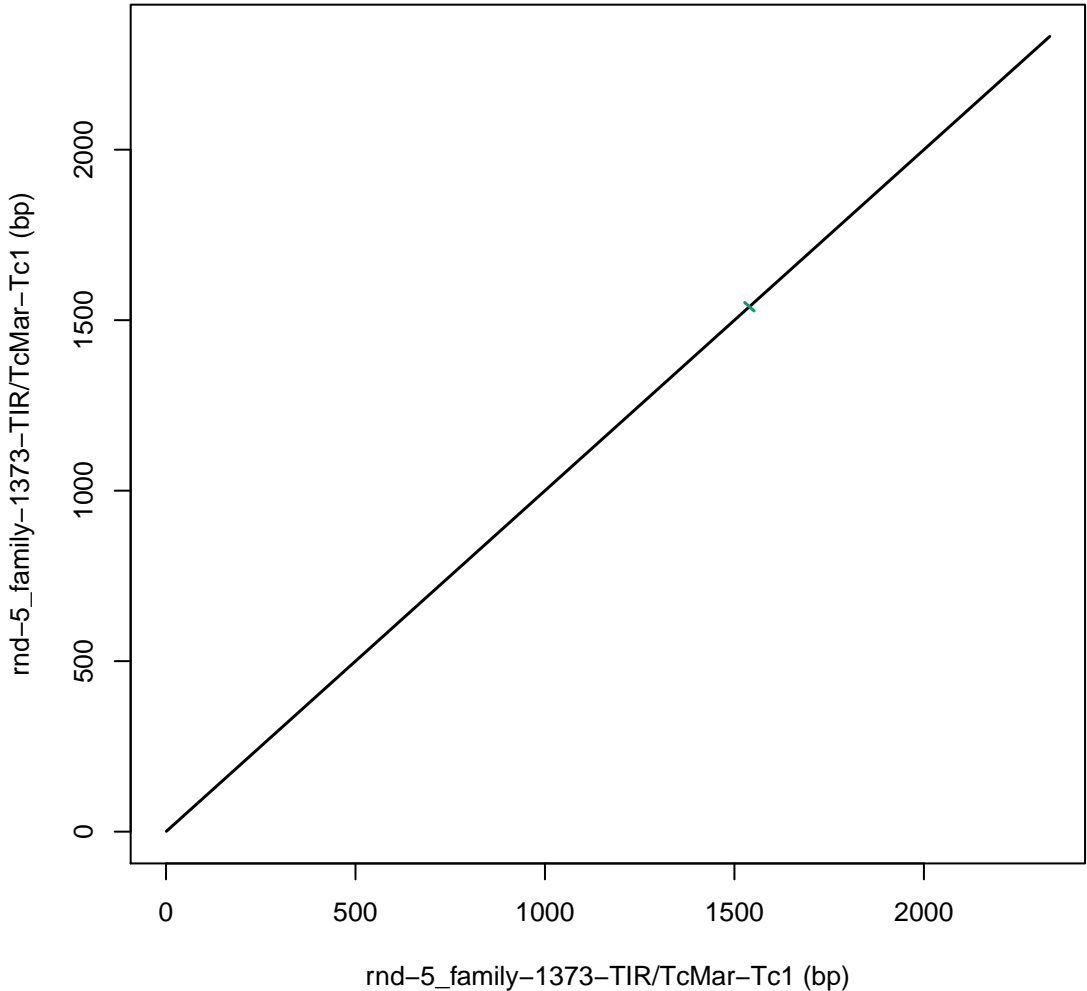
TE: rnd-5_family-1373-TIR/TcMar-Tc1
consensus size: 2332bp; fragments: 119; full length: 0 (>=2098.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

