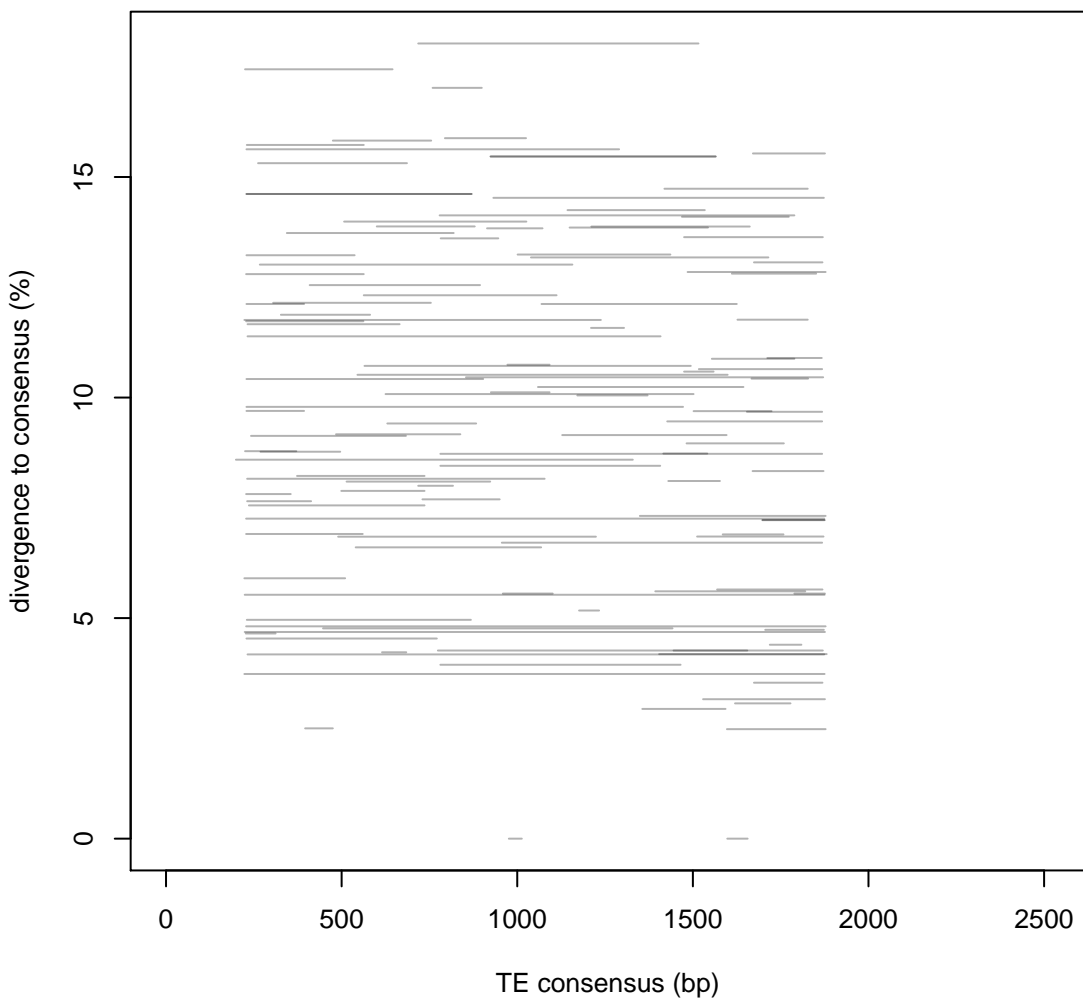
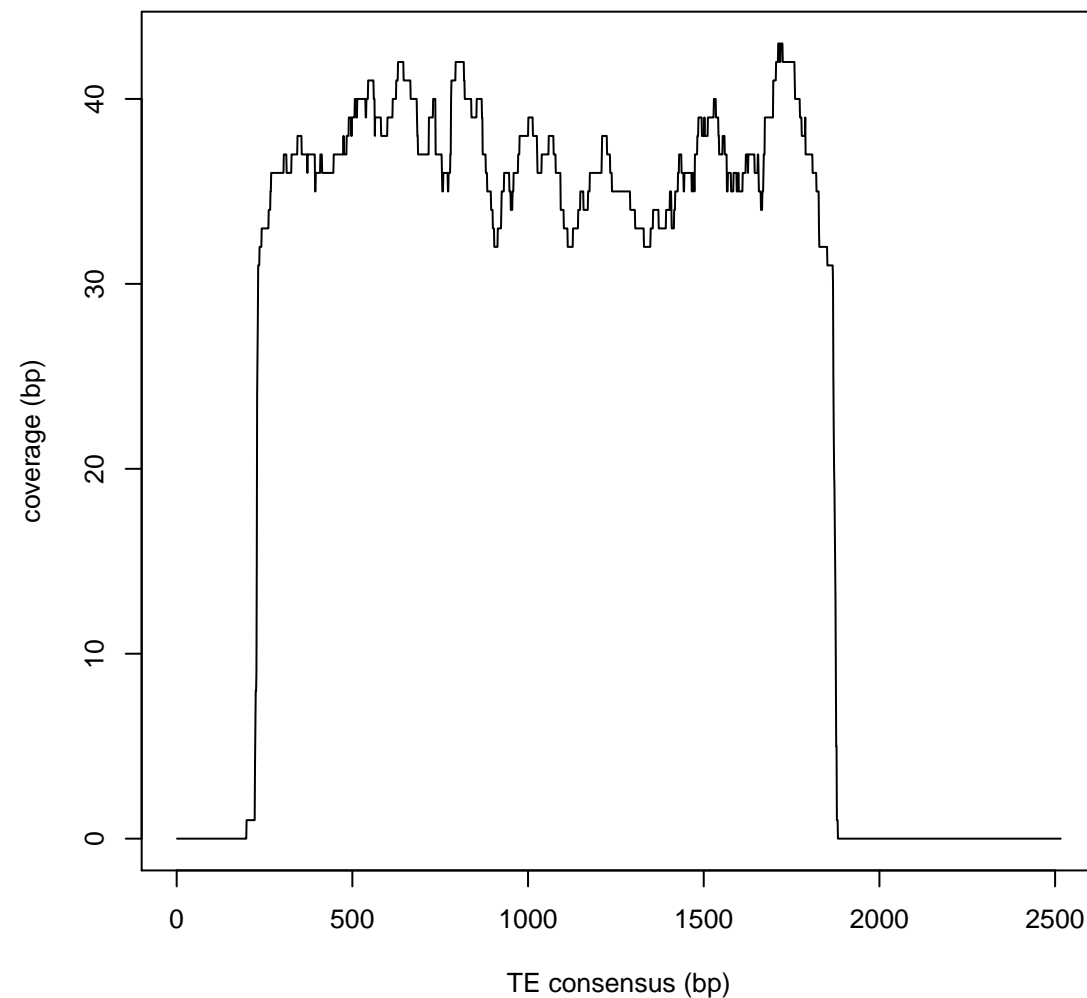


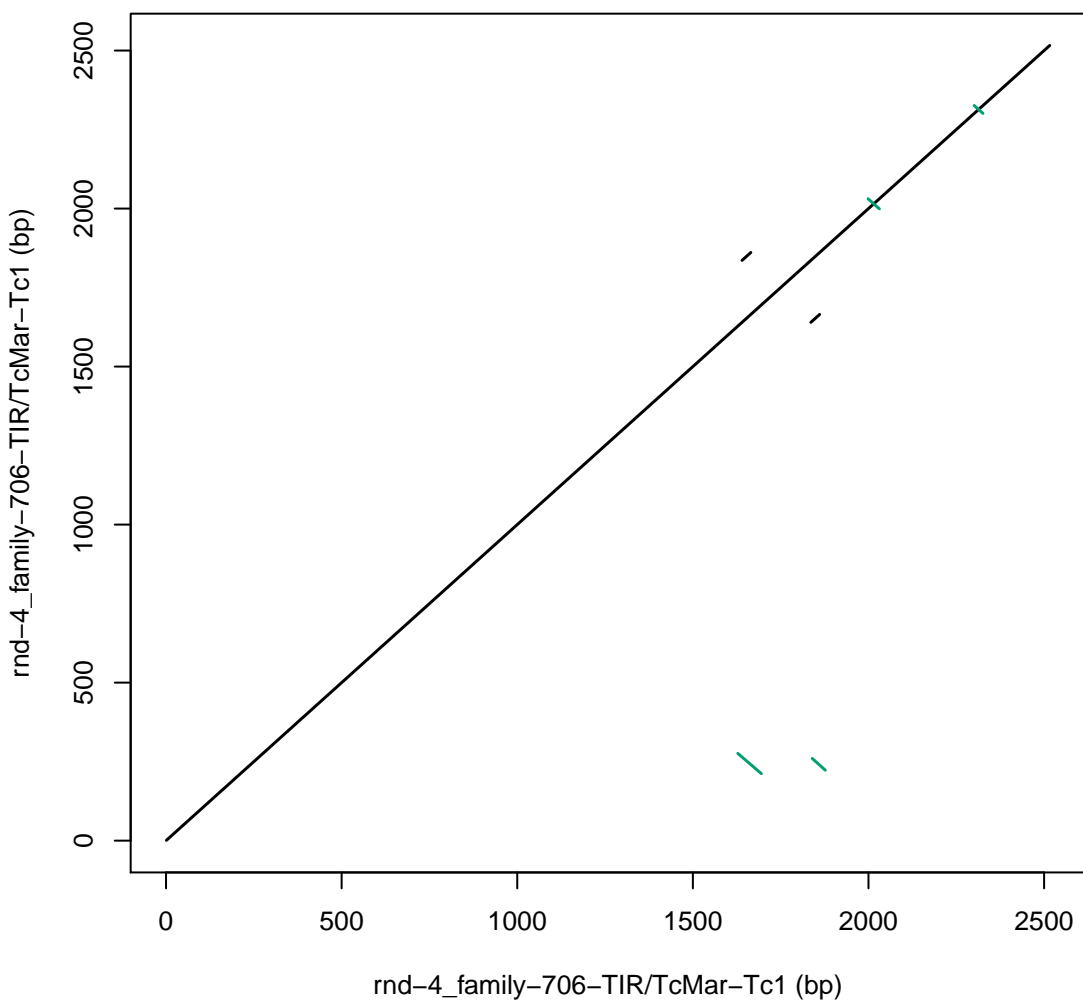
TE: rnd-4_family-706-TIR/TcMar-Tc1
consensus size: 2516bp; fragments: 127; full length: 0 (>=2264.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

