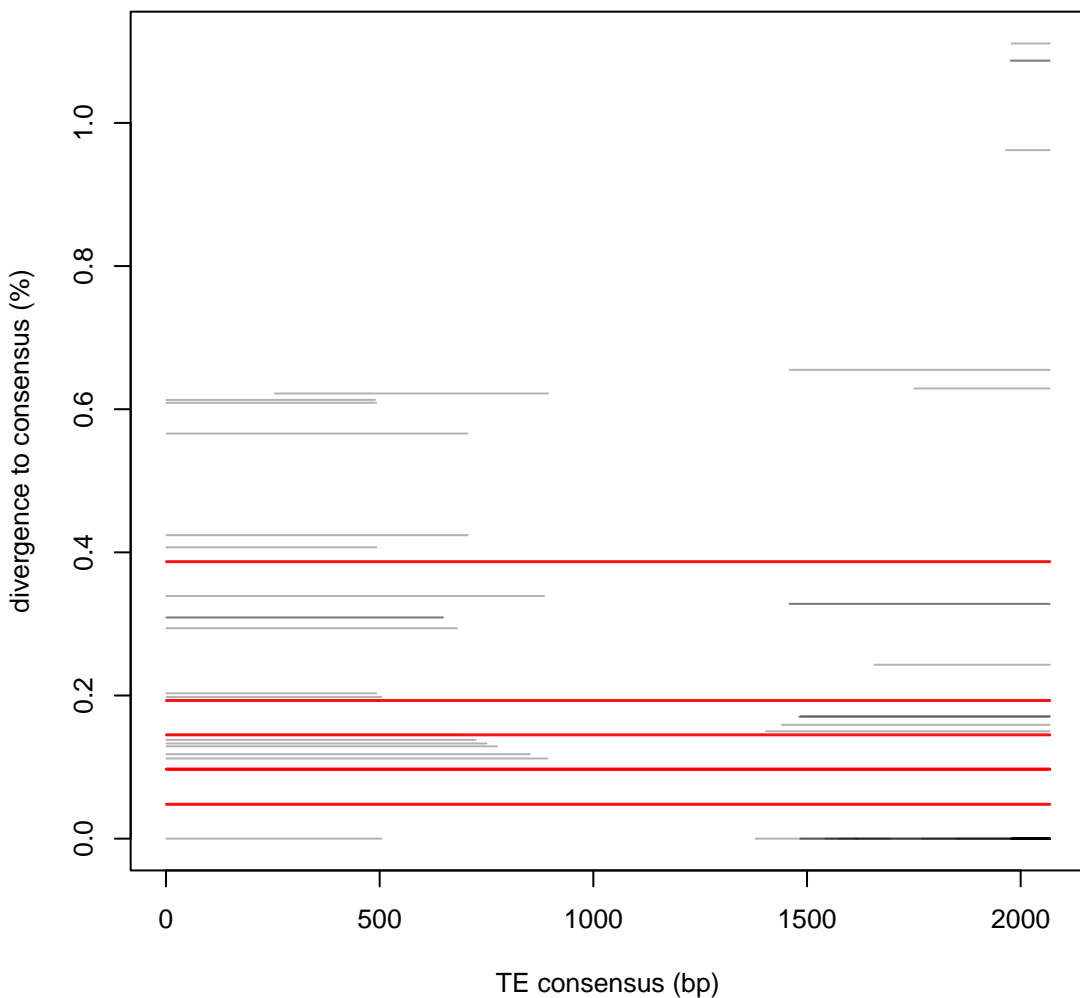
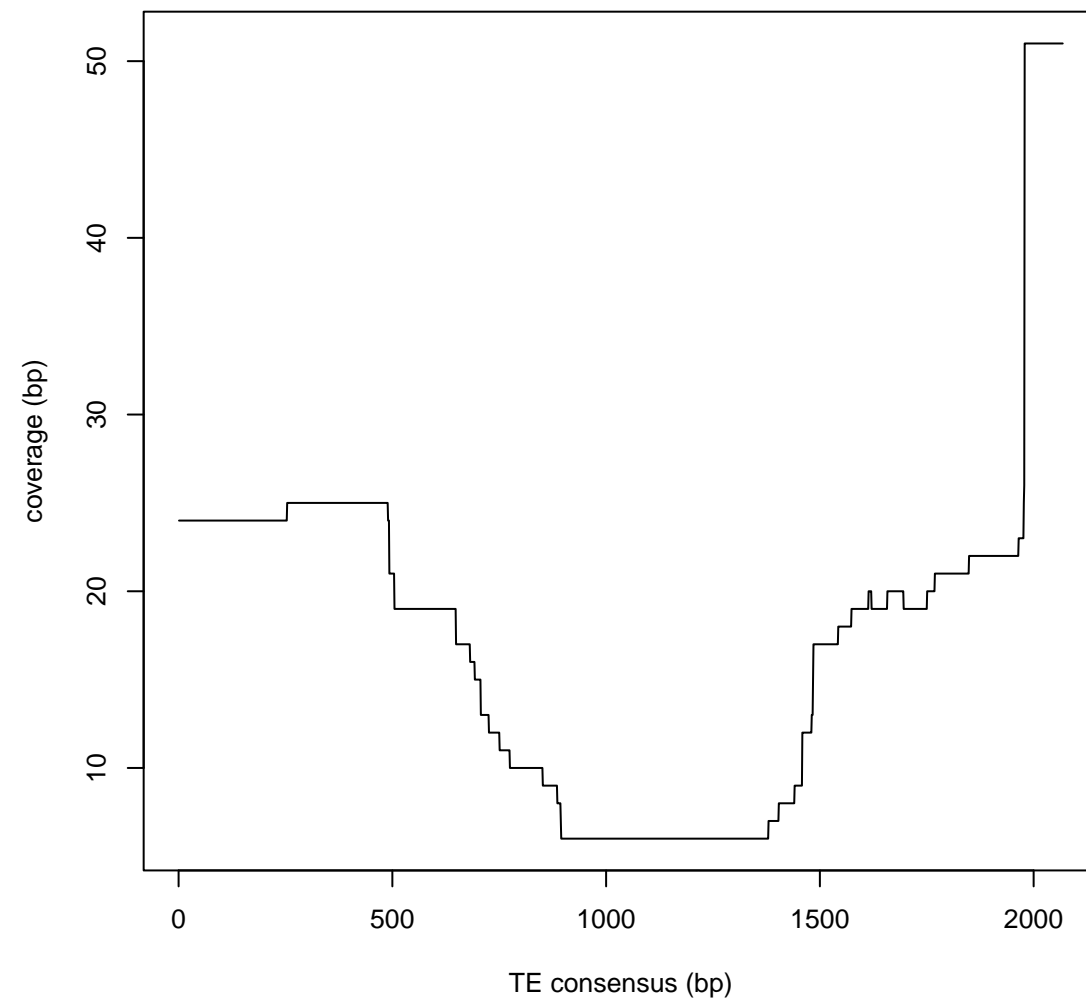


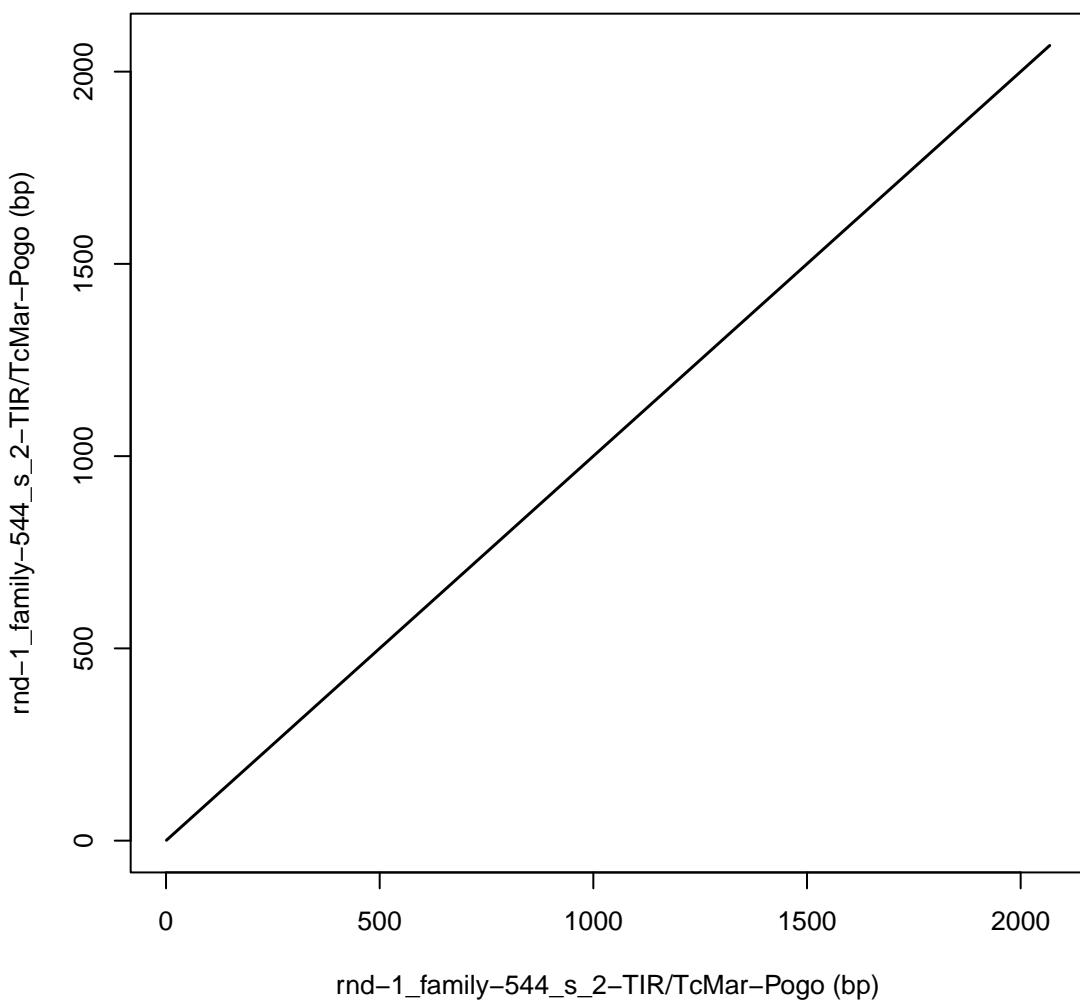
TE: rnd-1_family-544_s_2-TIR/TcMar-Pogo
consensus size: 2068bp; fragments: 72; full length: 6 (≥ 1861.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

