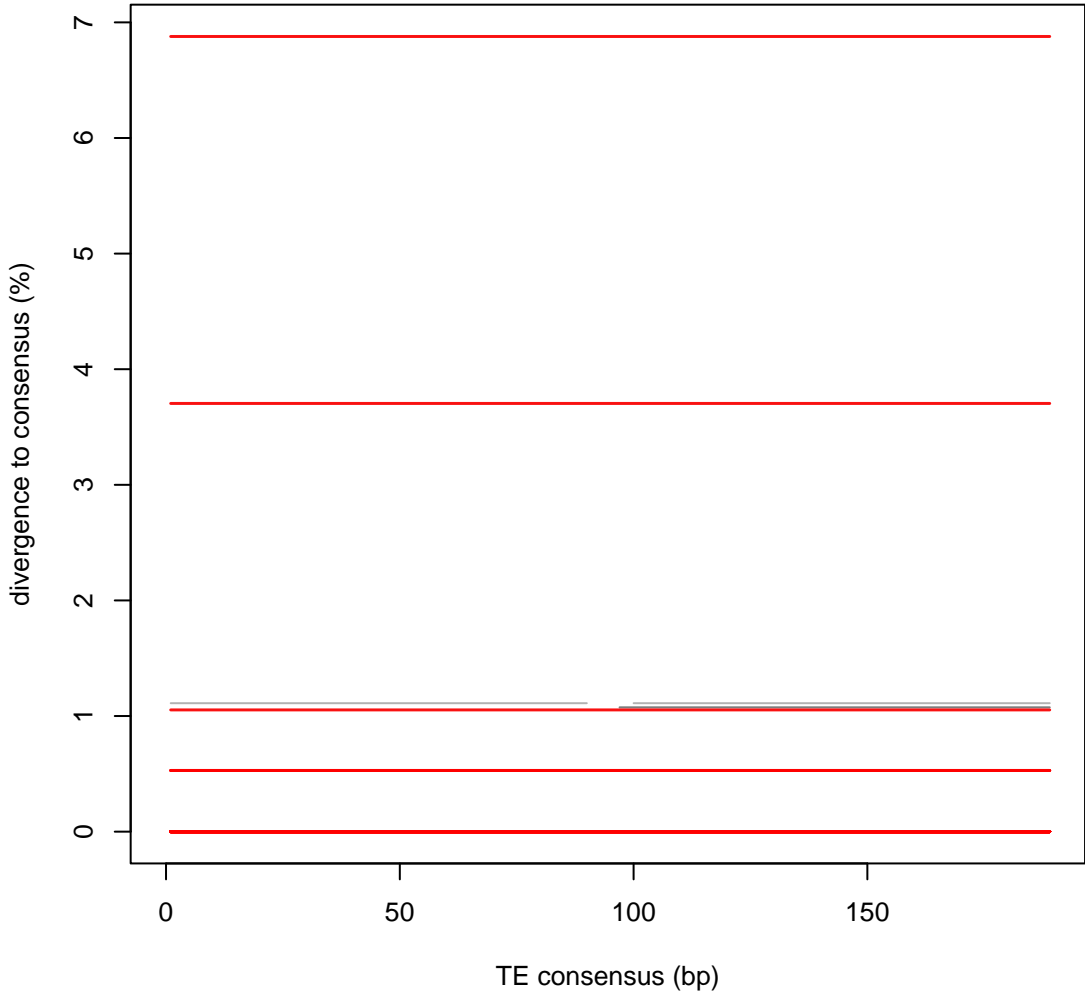
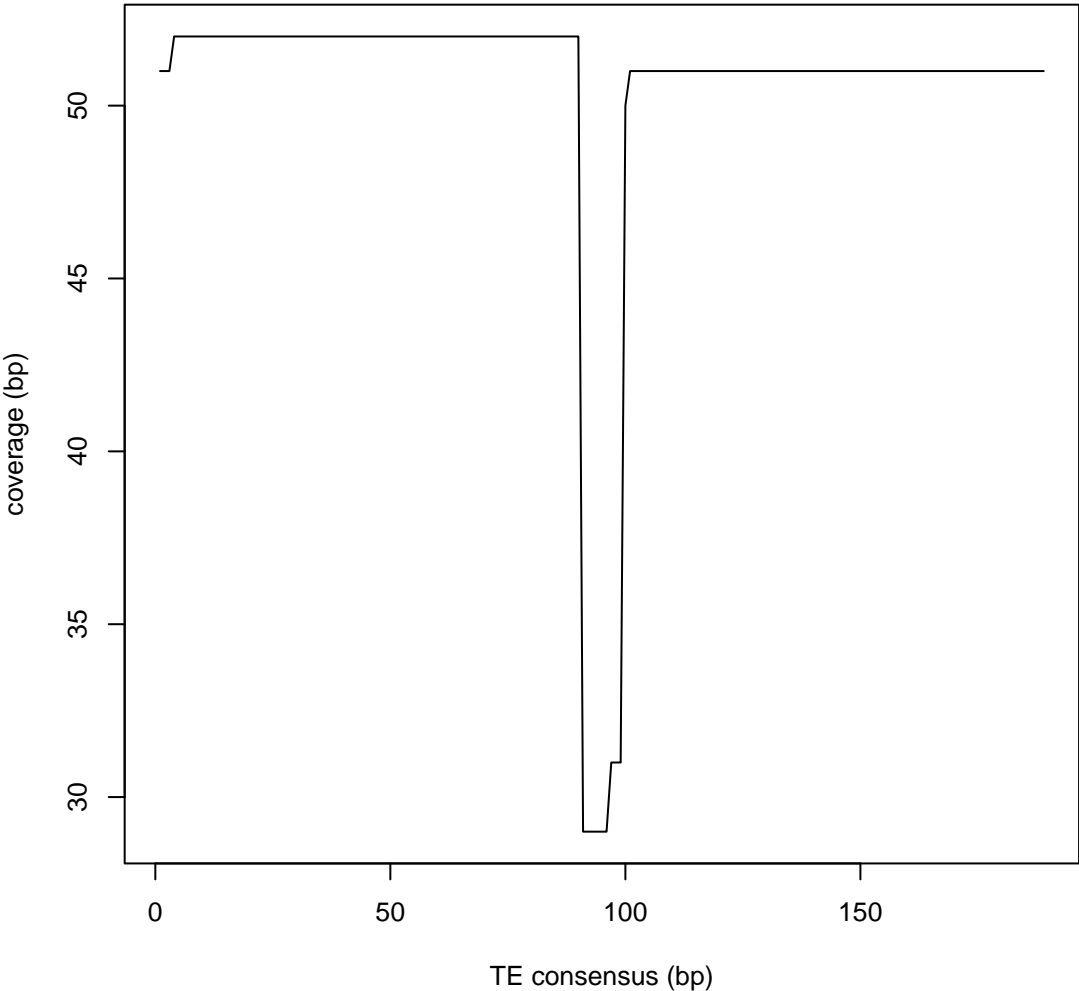


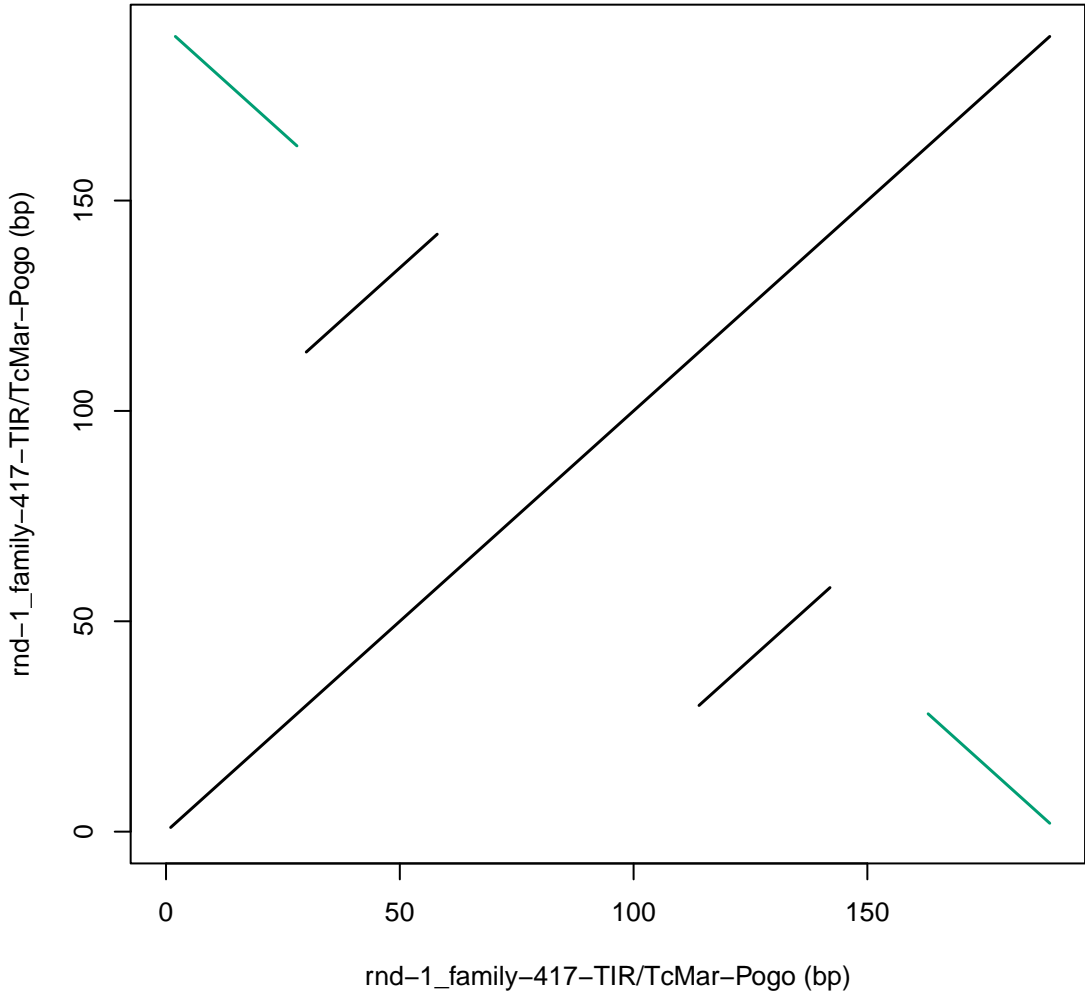
TE: rnd-1_family-417-TIR/TcMar-Pogo
consensus size: 189bp; fragments: 74; full length: 29 (>=170.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

