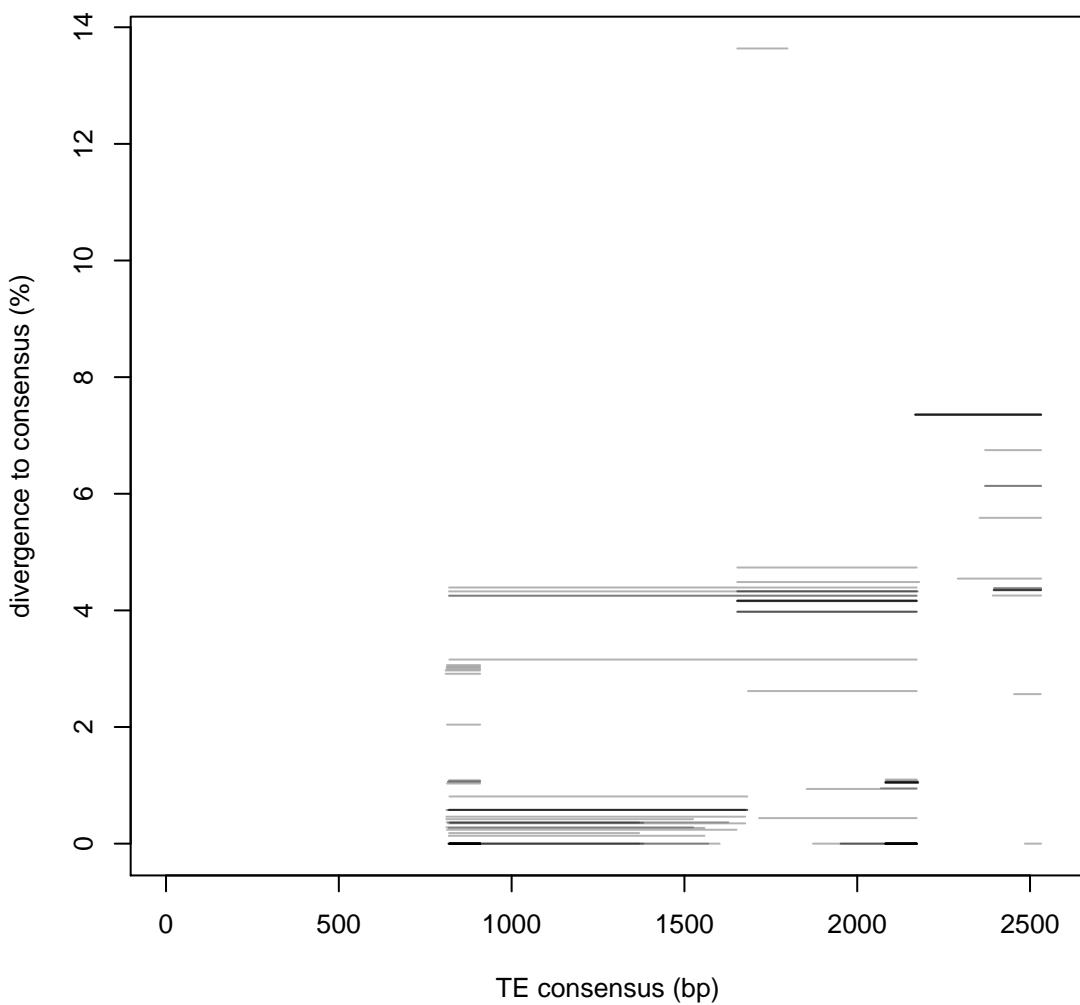
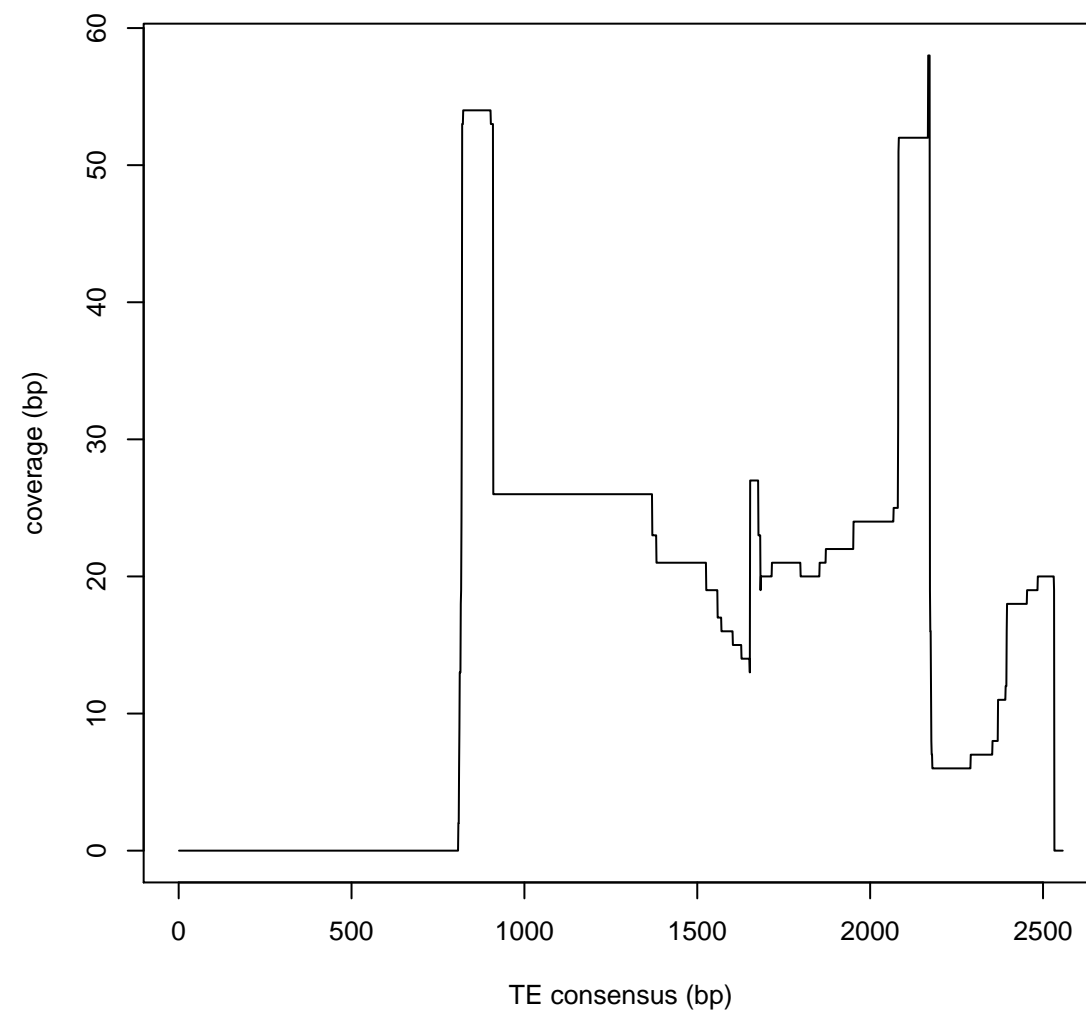


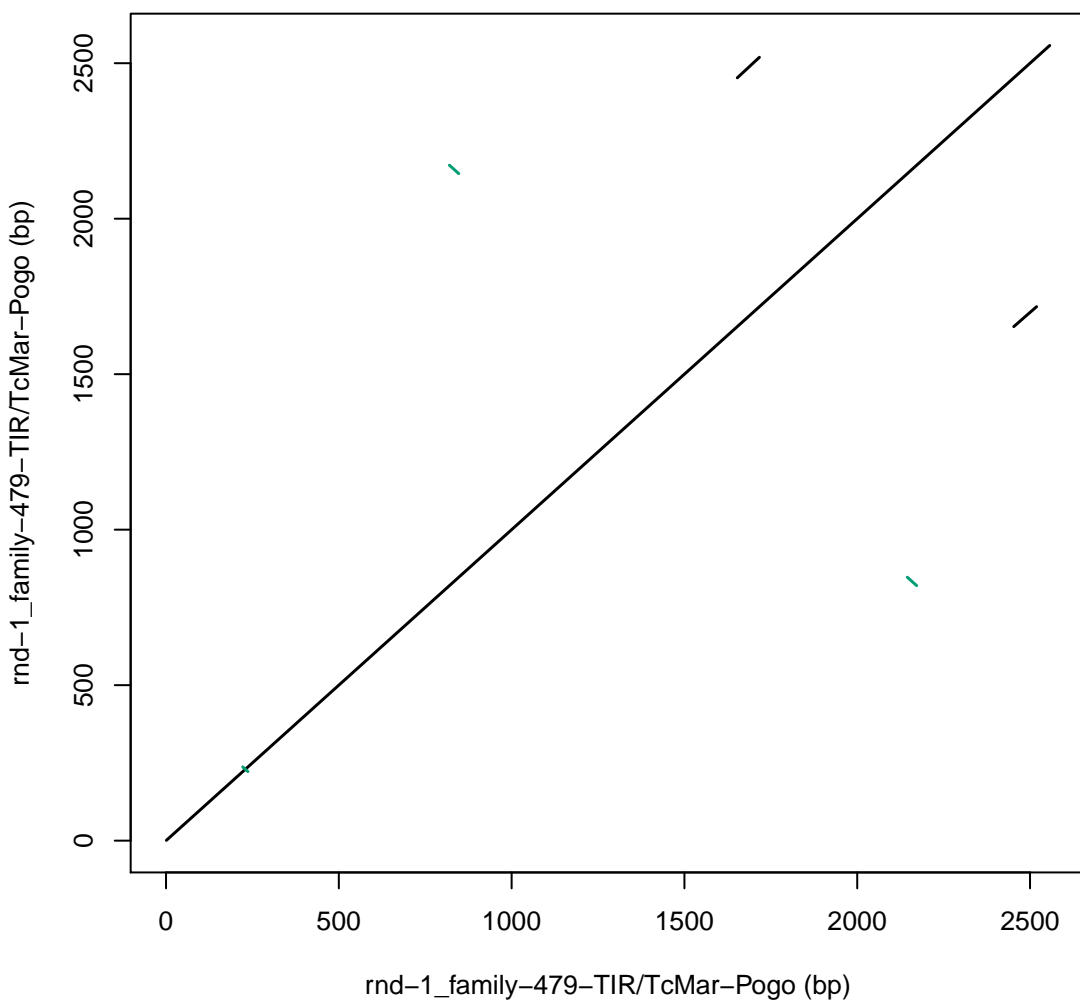
TE: rnd-1_family-479-TIR/TcMar-Pogo
consensus size: 2557bp; fragments: 122; full length: 0 (>=2301.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

