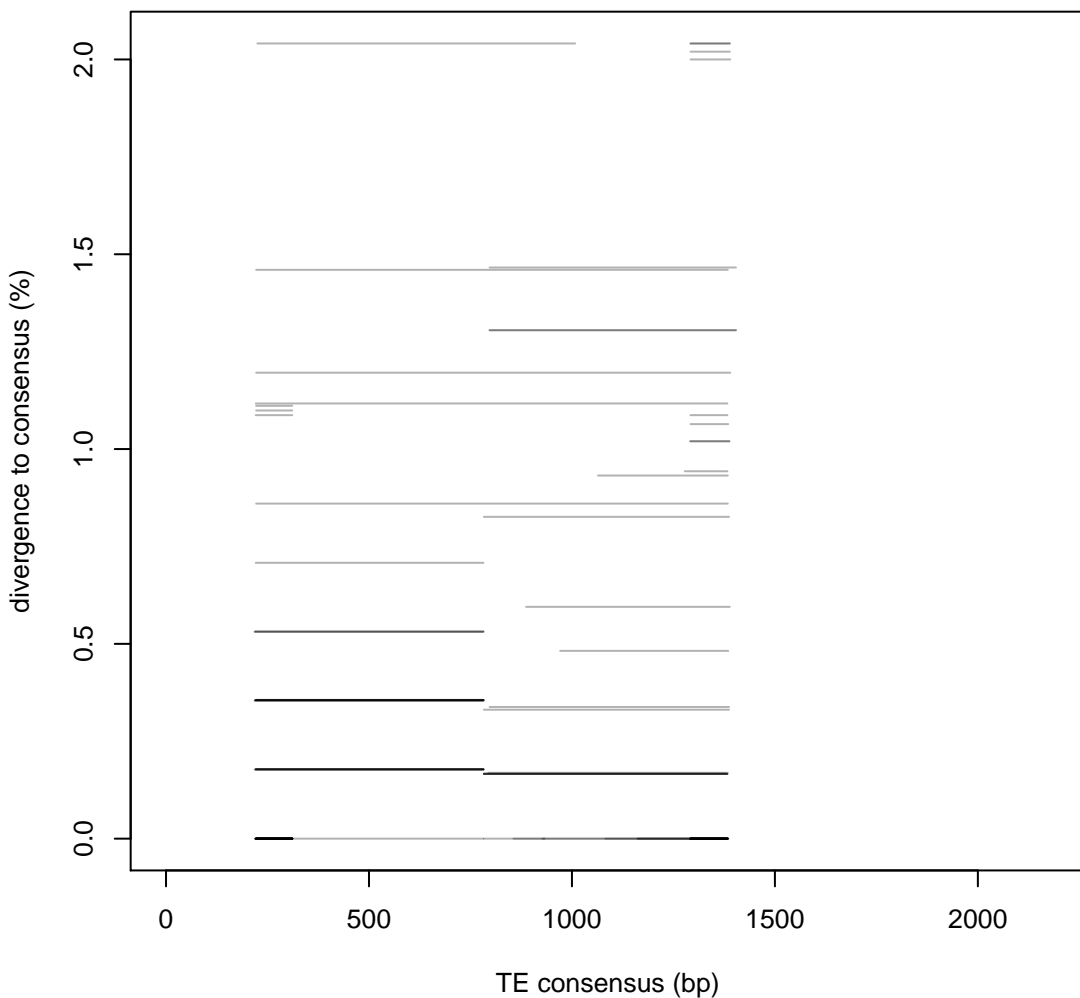
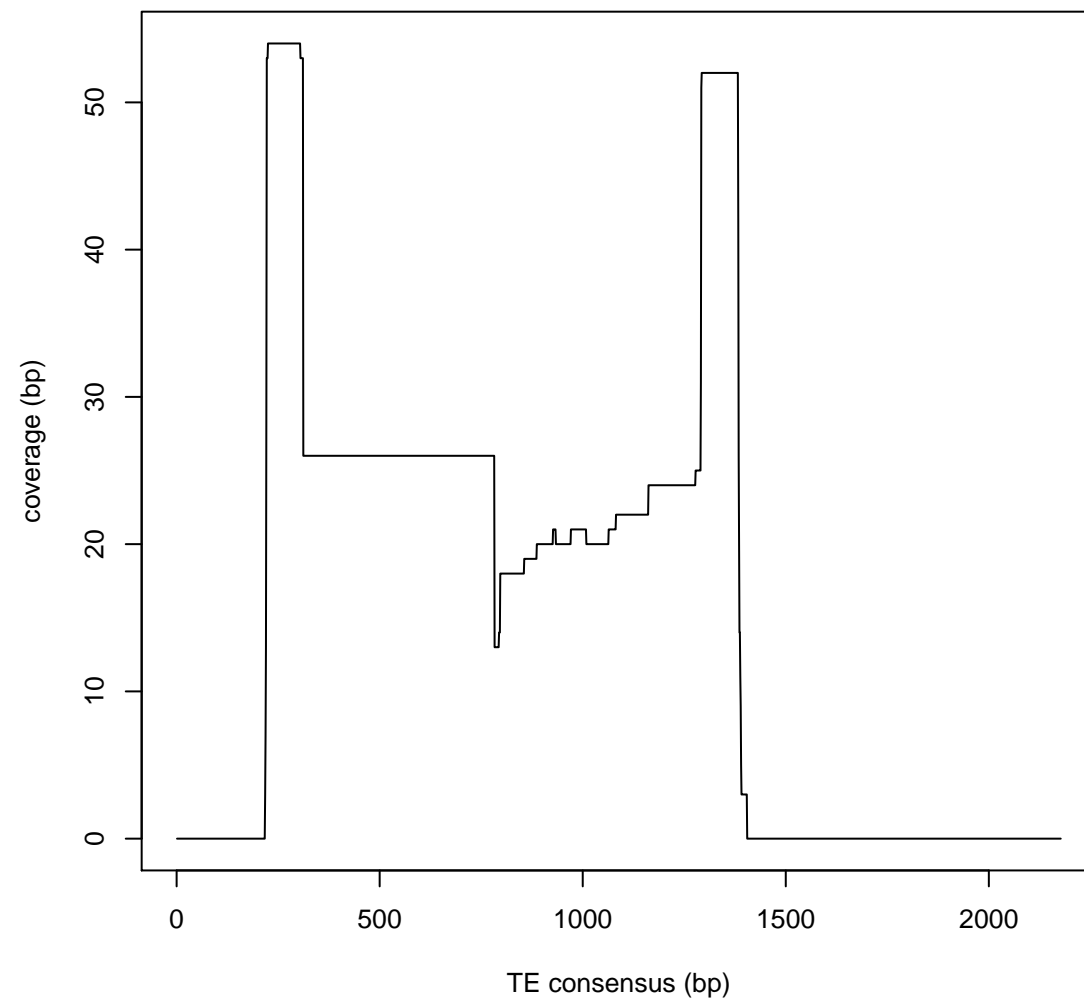


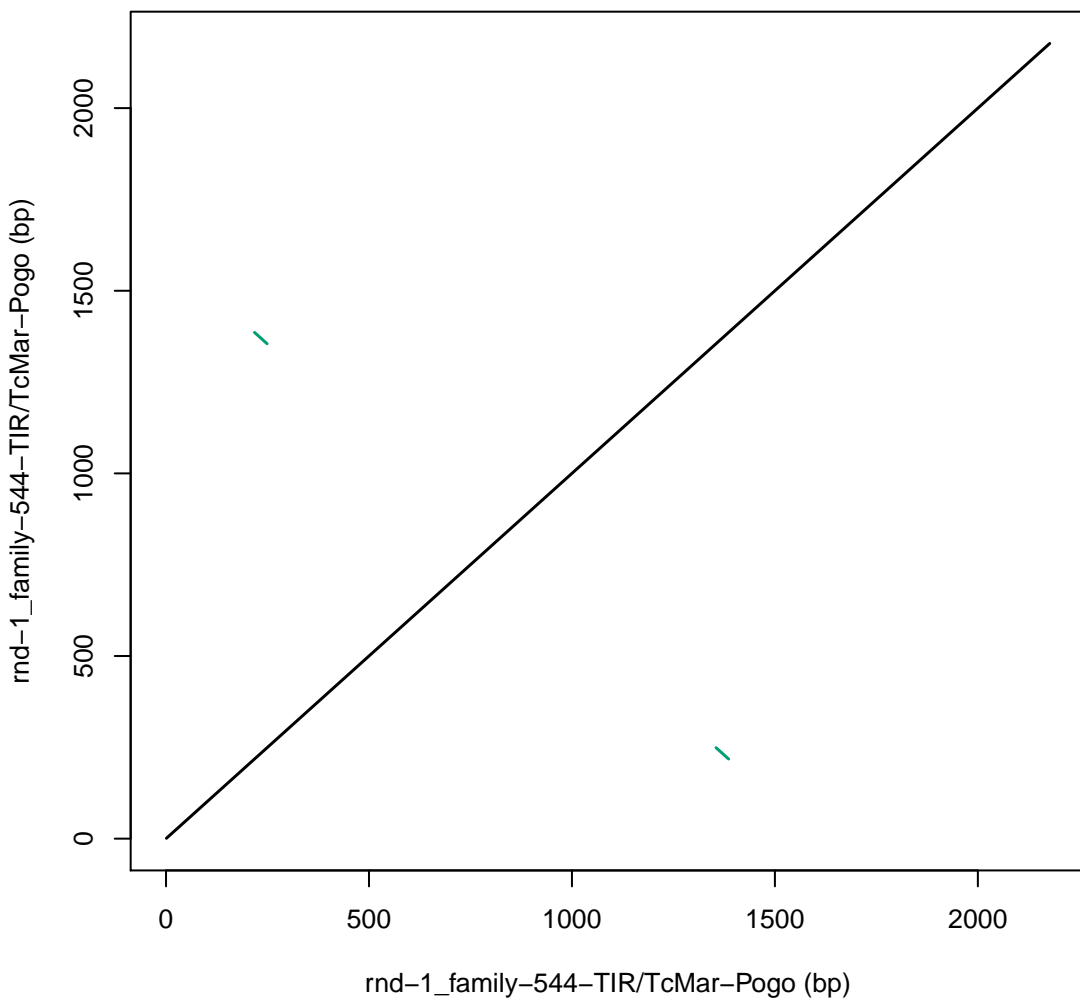
TE: rnd-1\_family-544-TIR/TcMar-Pogo  
consensus size: 2177bp; fragments: 103; full length: 0 (>=1959.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

