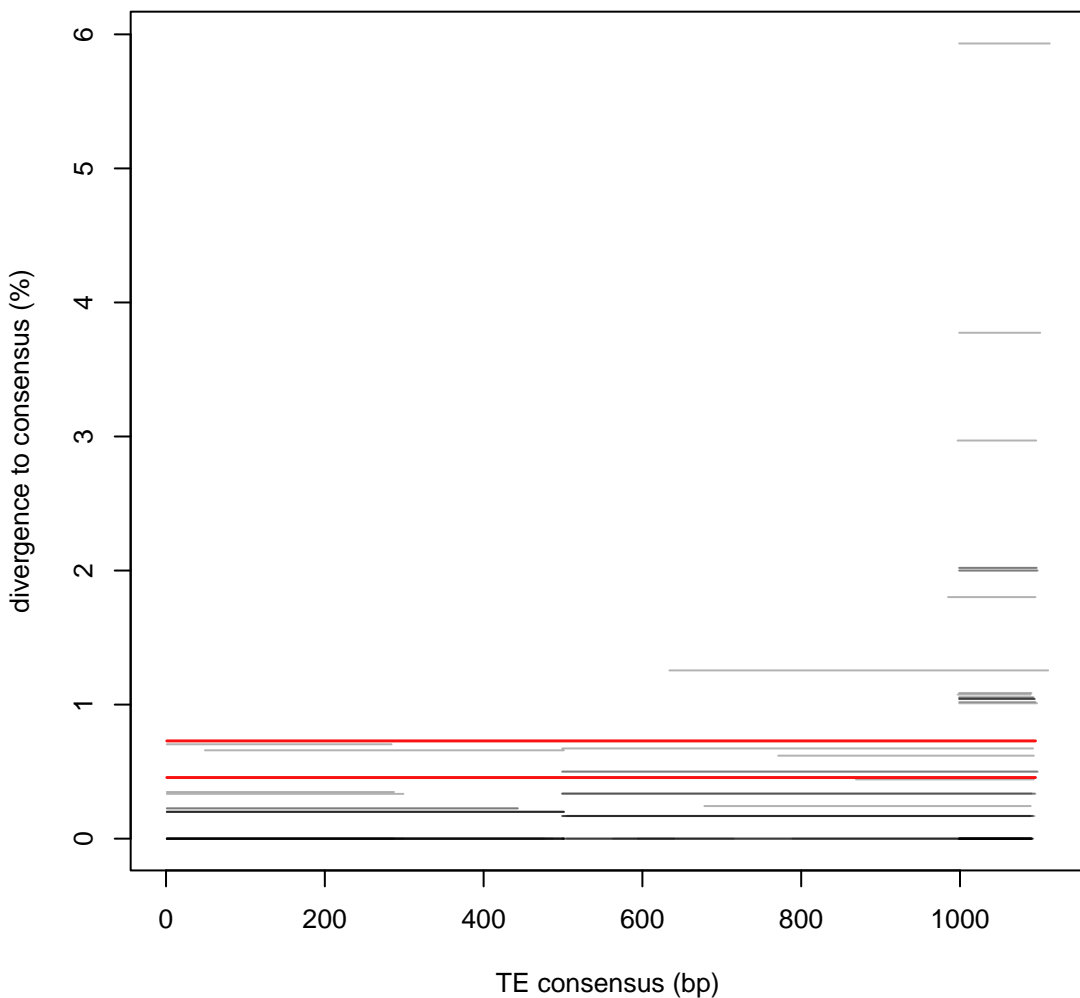
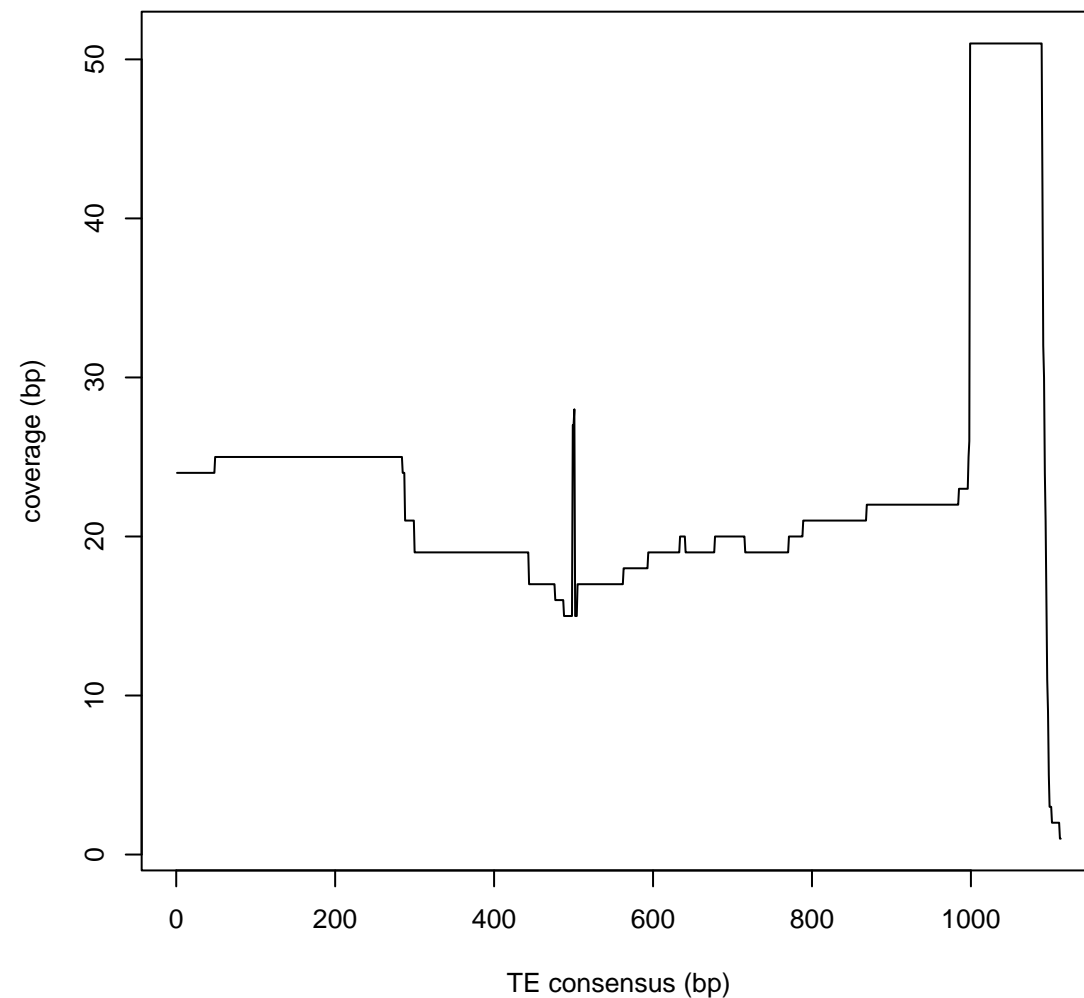


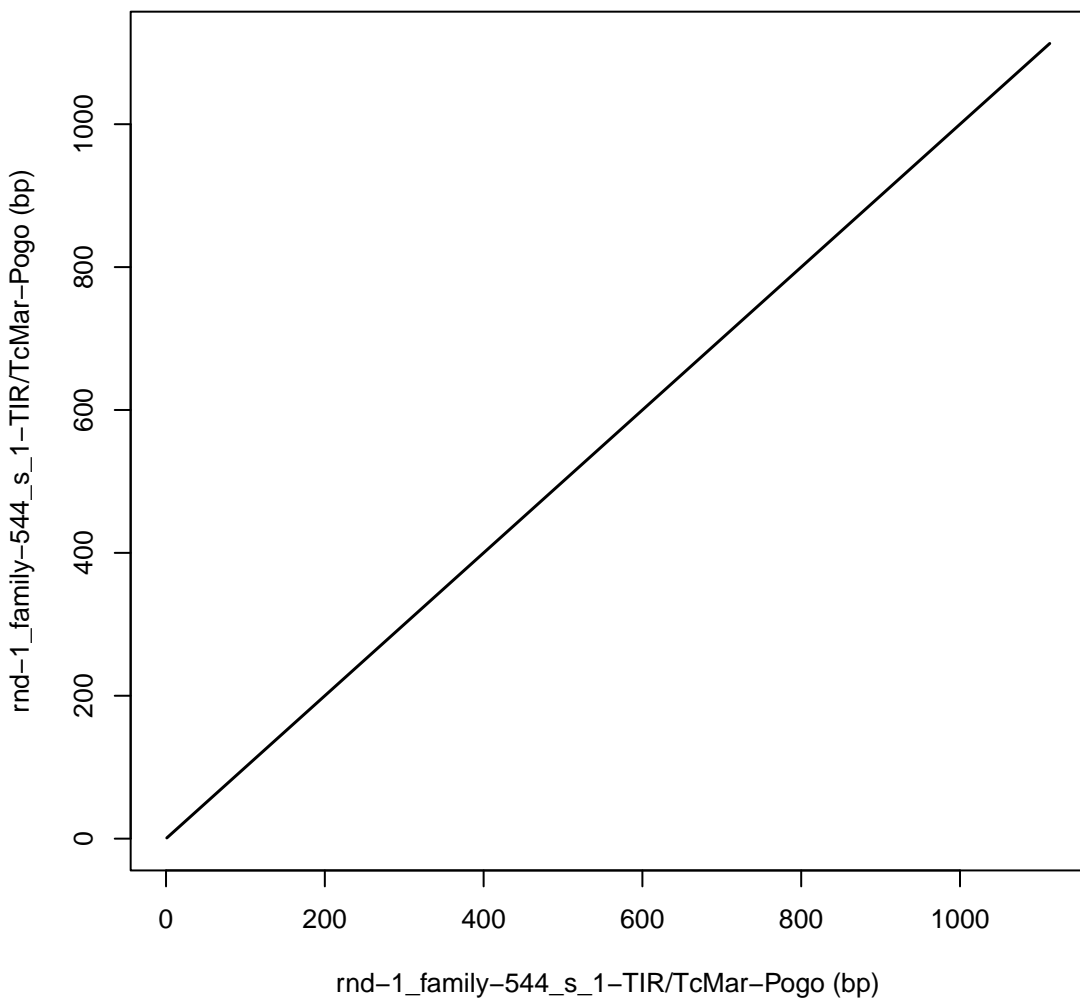
TE: rnd-1\_family-544\_s\_1-TIR/TcMar-Pogo  
consensus size: 1113bp; fragments: 76; full length: 2 ( $\geq 1001.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

