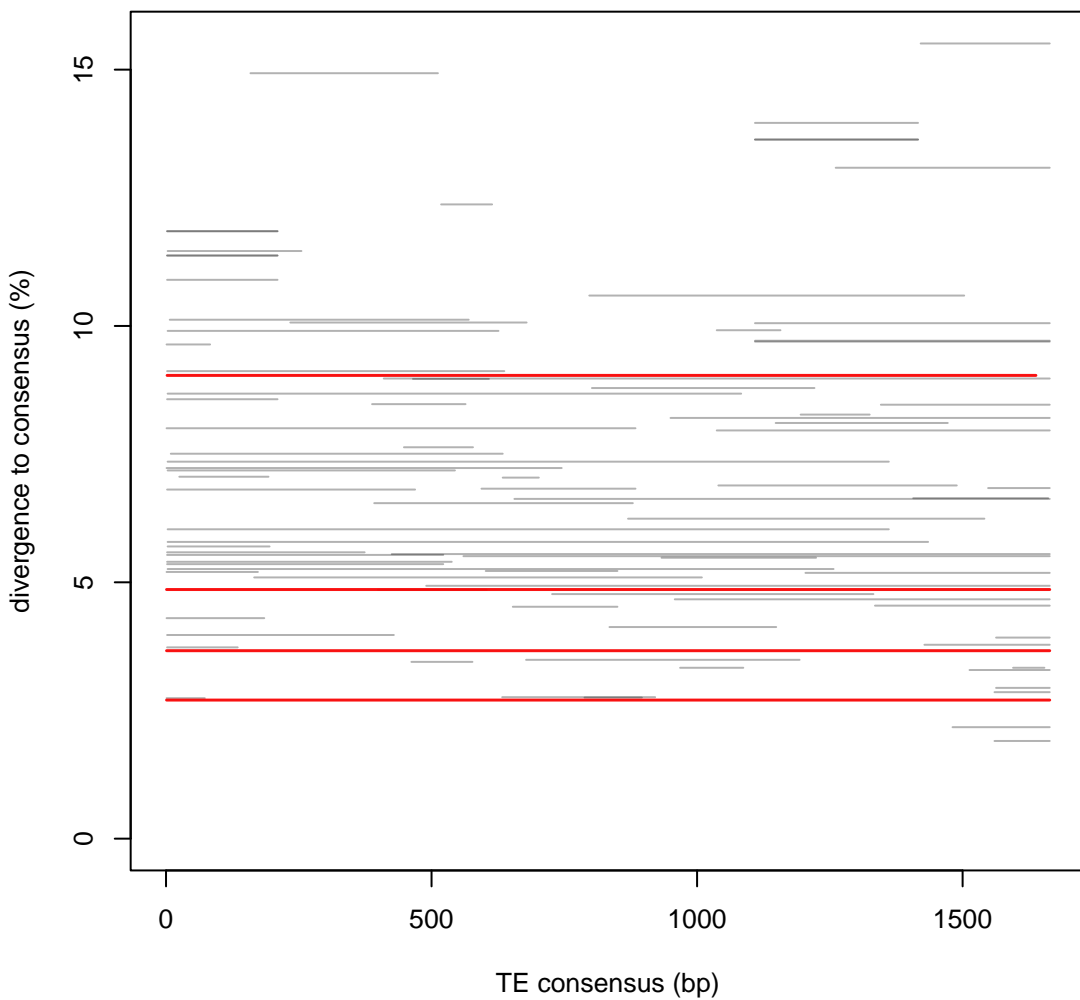
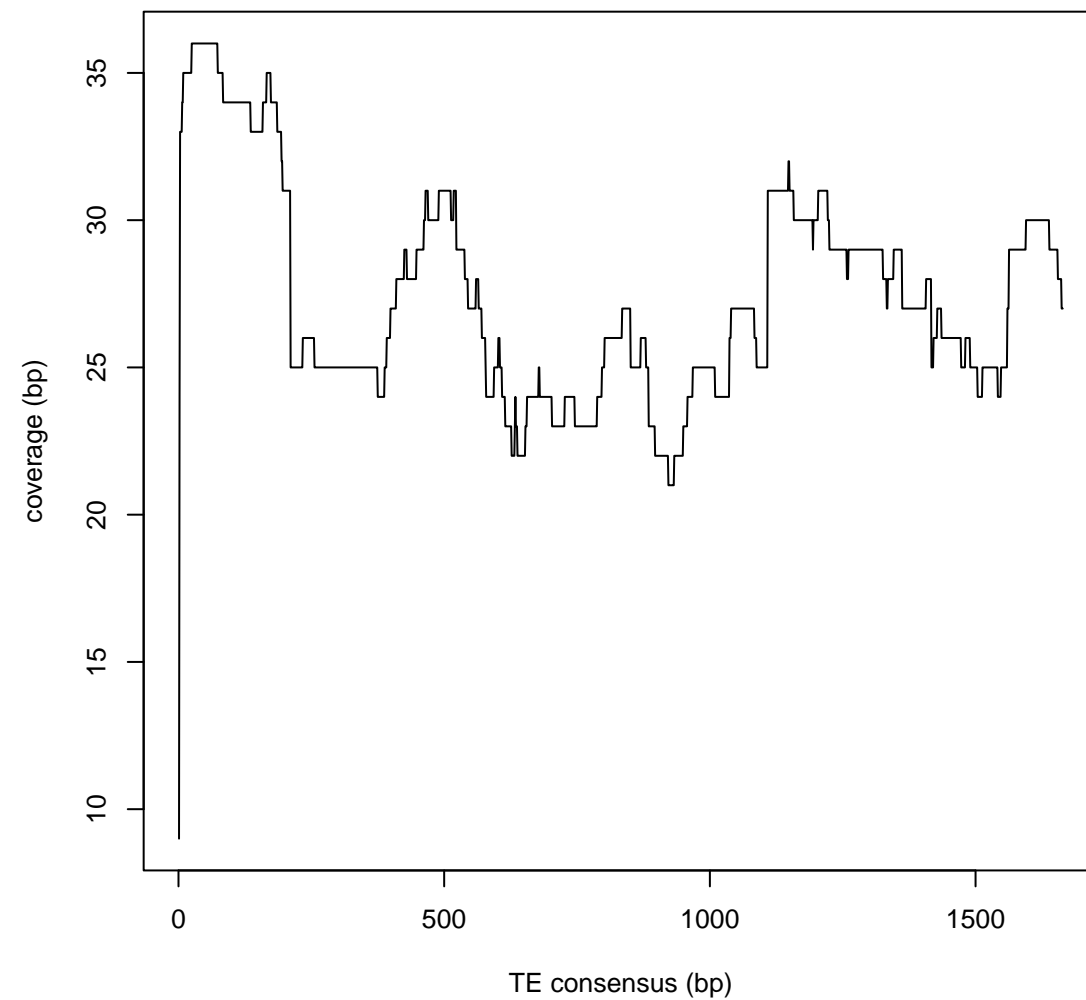


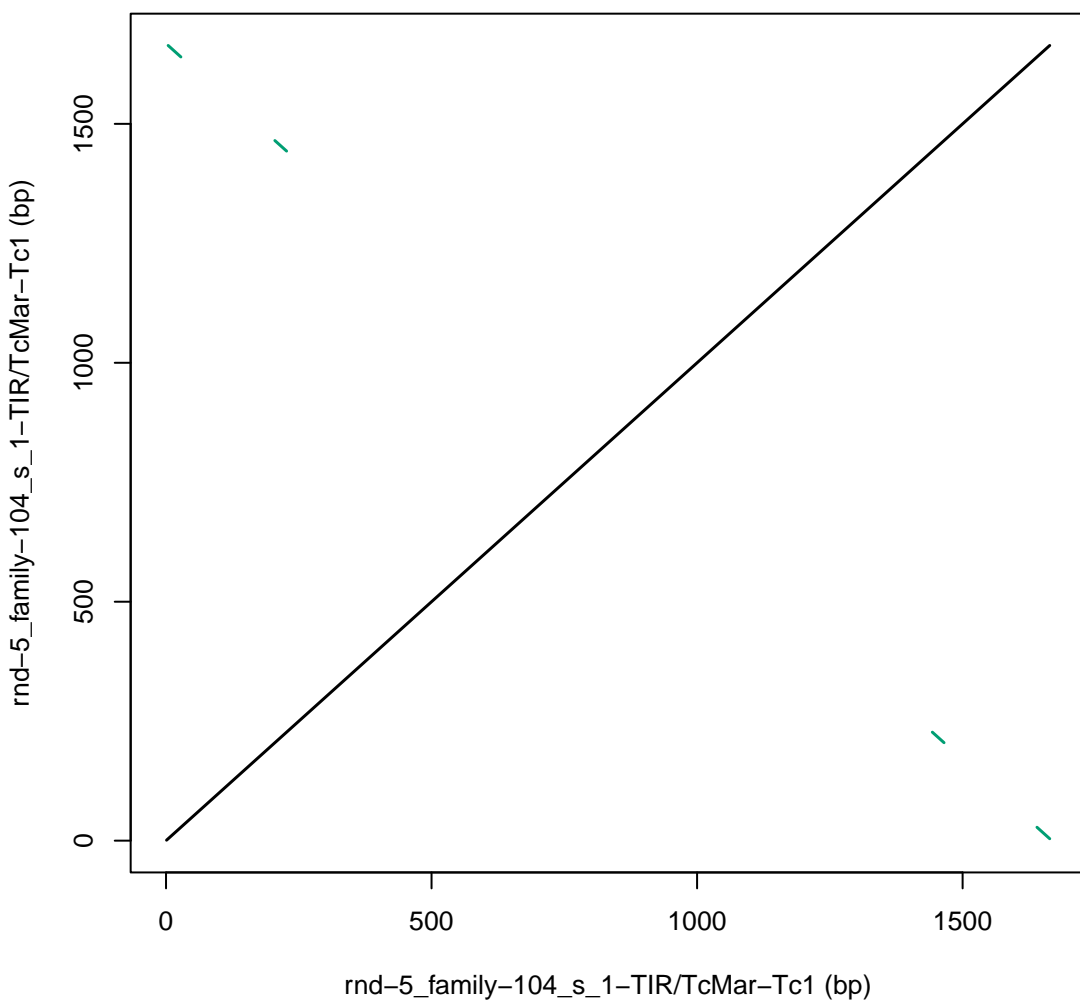
TE: rnd-5\_family-104\_s\_1-TIR/TcMar-Tc1  
consensus size: 1664bp; fragments: 93; full length: 4 ( $\geq 1497.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

