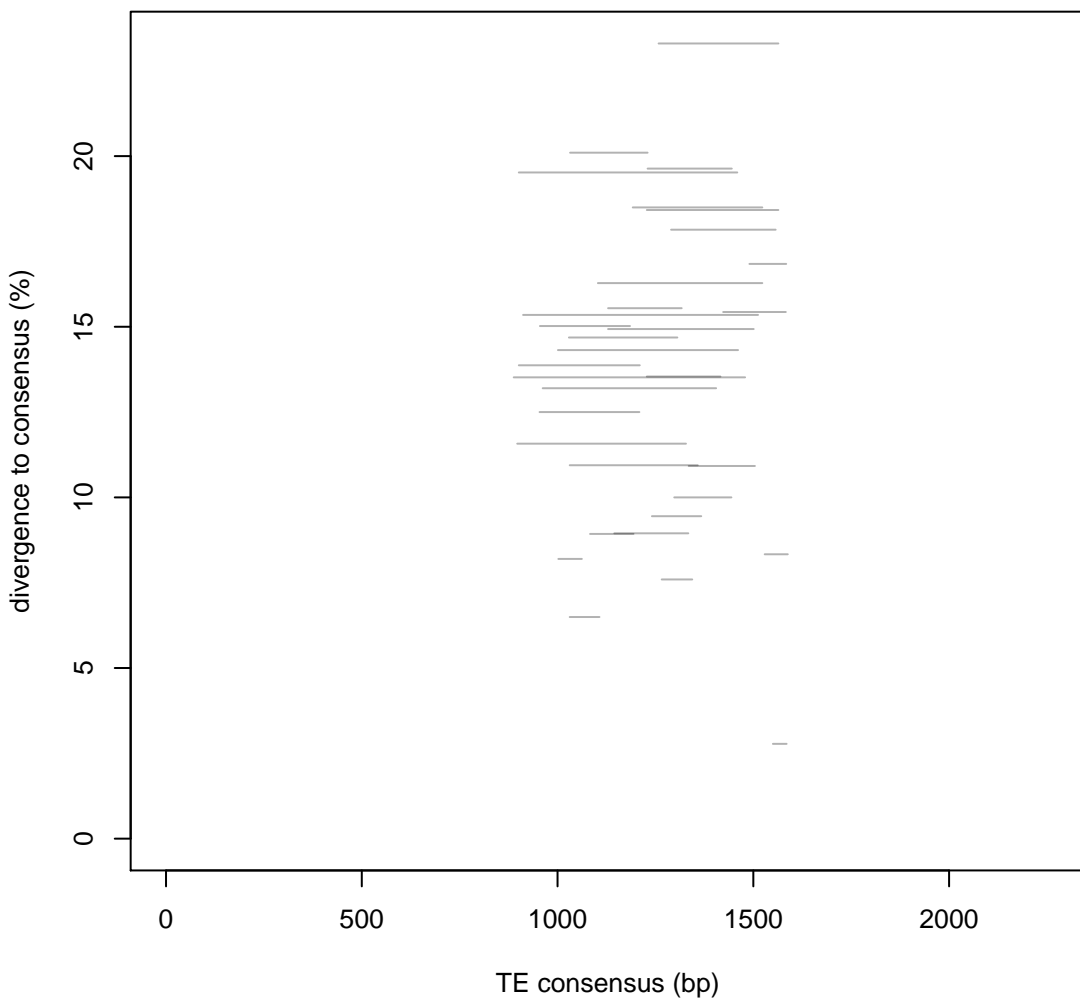
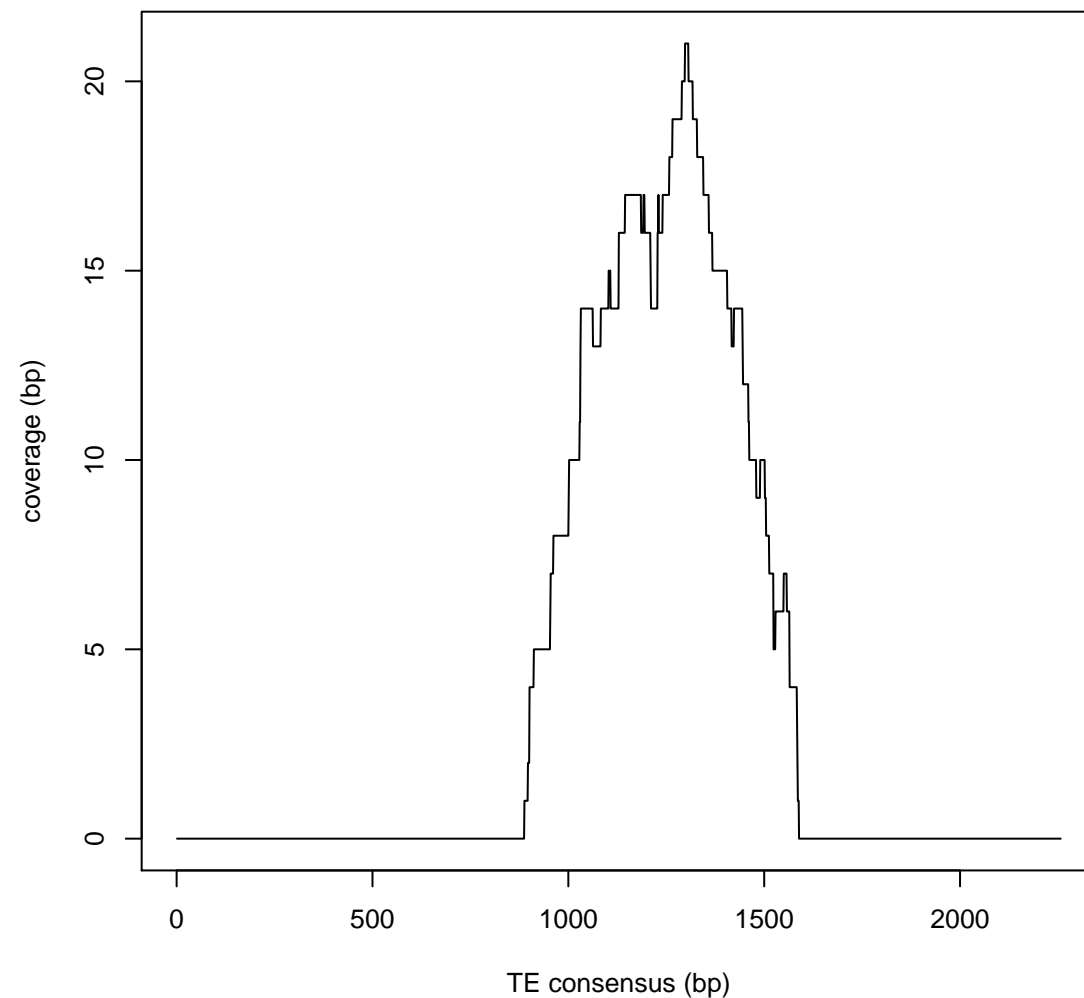


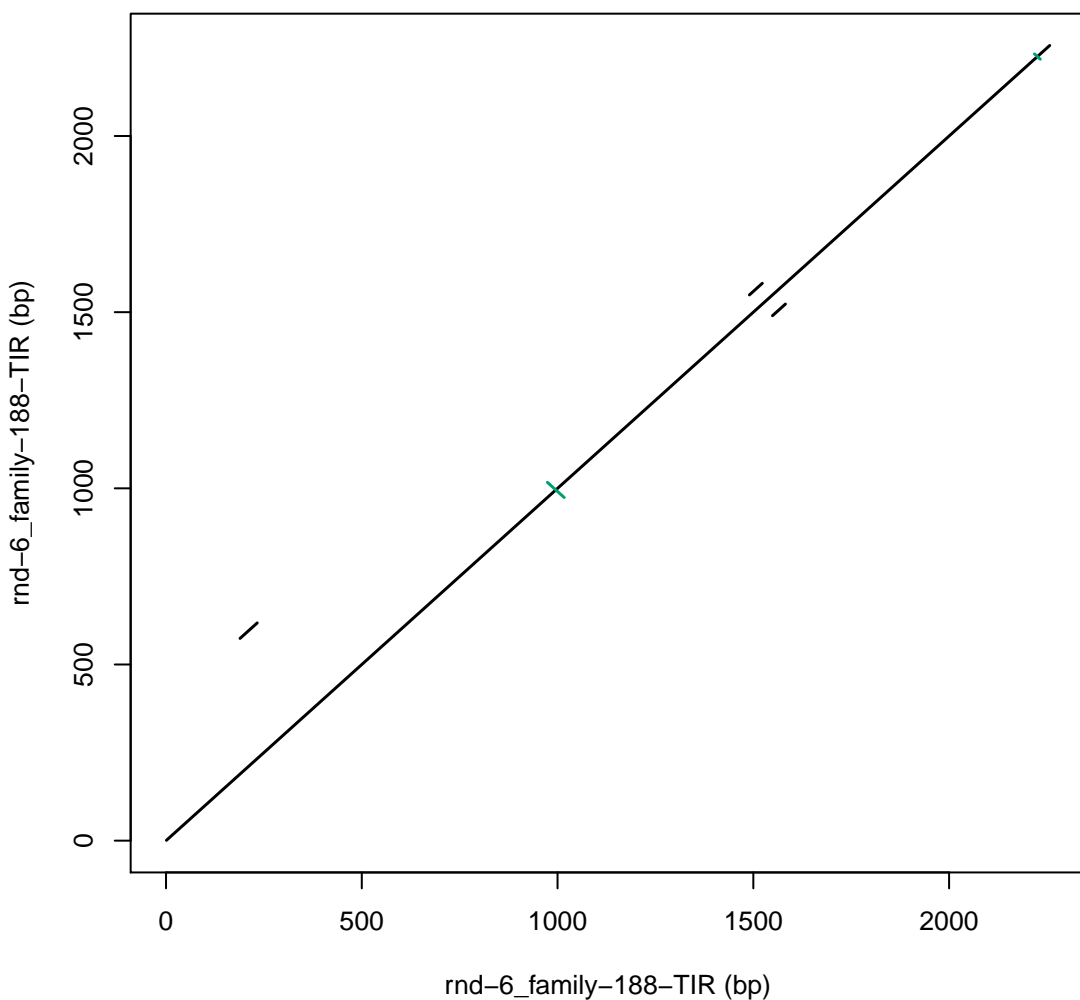
TE: rnd-6\_family-188-TIR  
consensus size: 2257bp; fragments: 33; full length: 0 ( $\geq 2031.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

