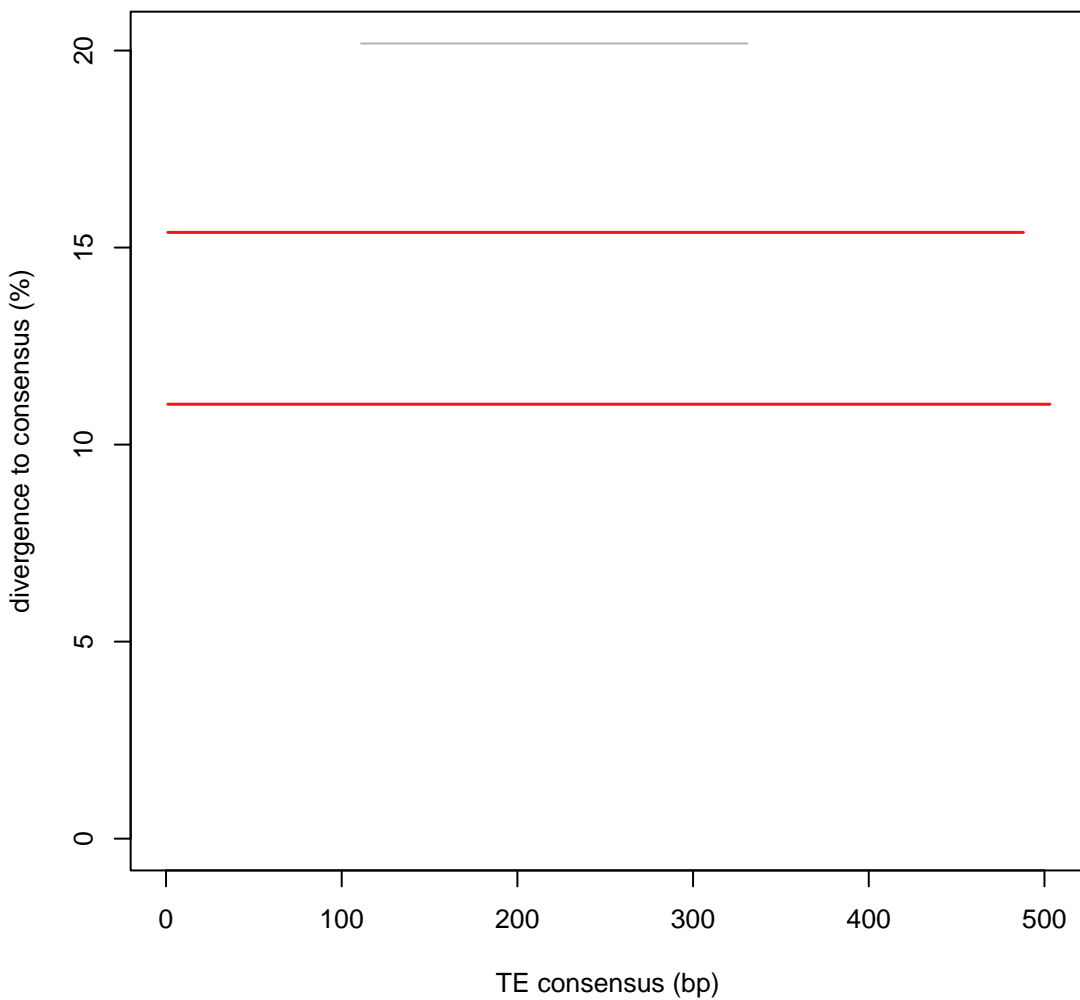
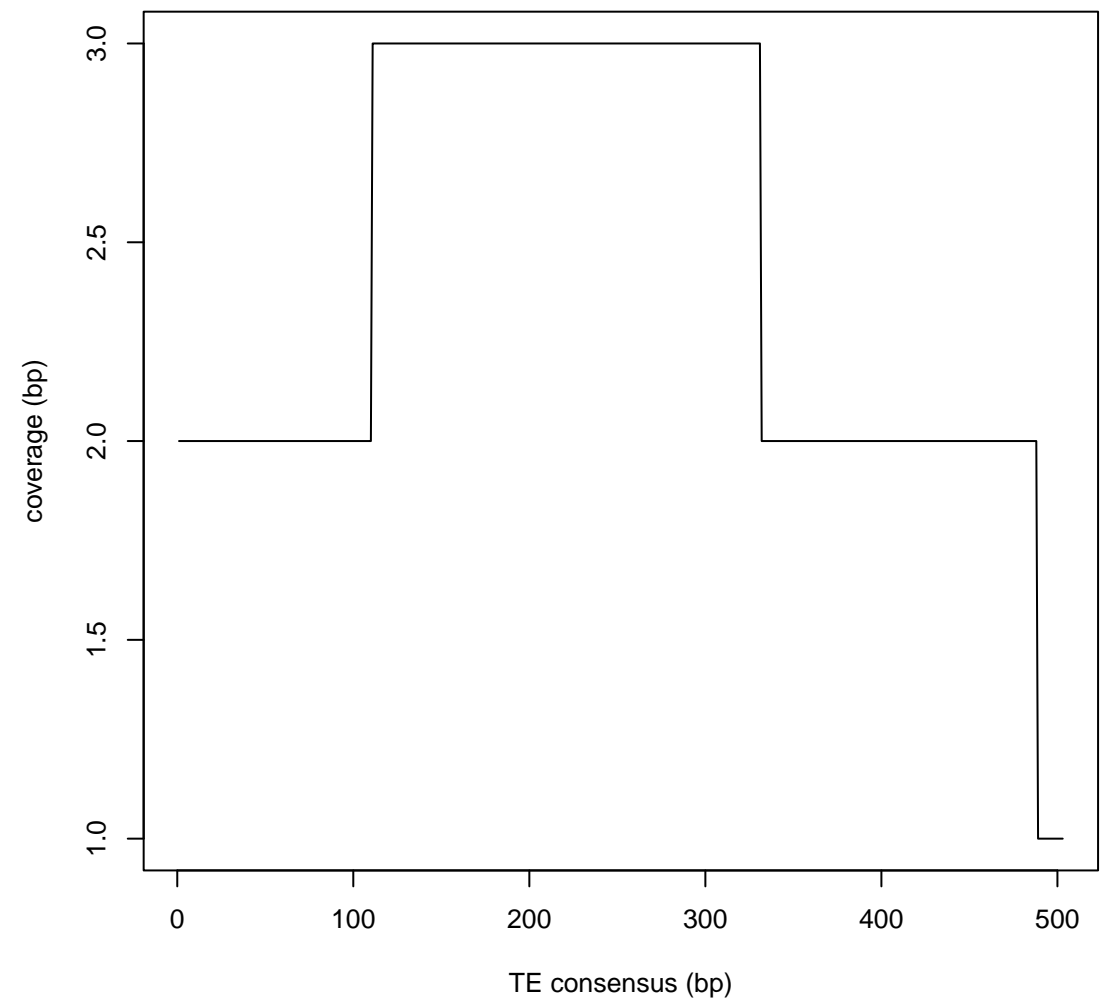


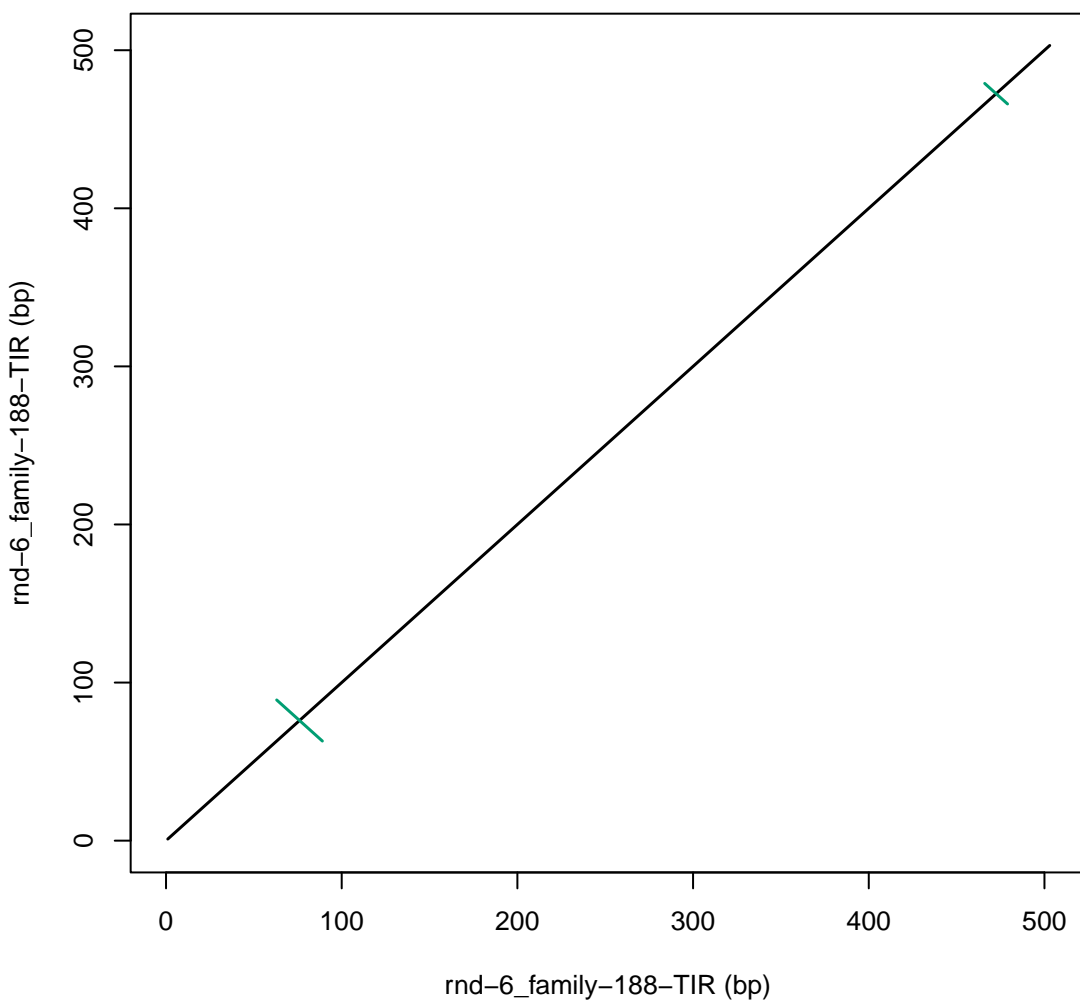
TE: rnd-6\_family-188-TIR  
consensus size: 503bp; fragments: 3; full length: 2 ( $\geq 452.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

