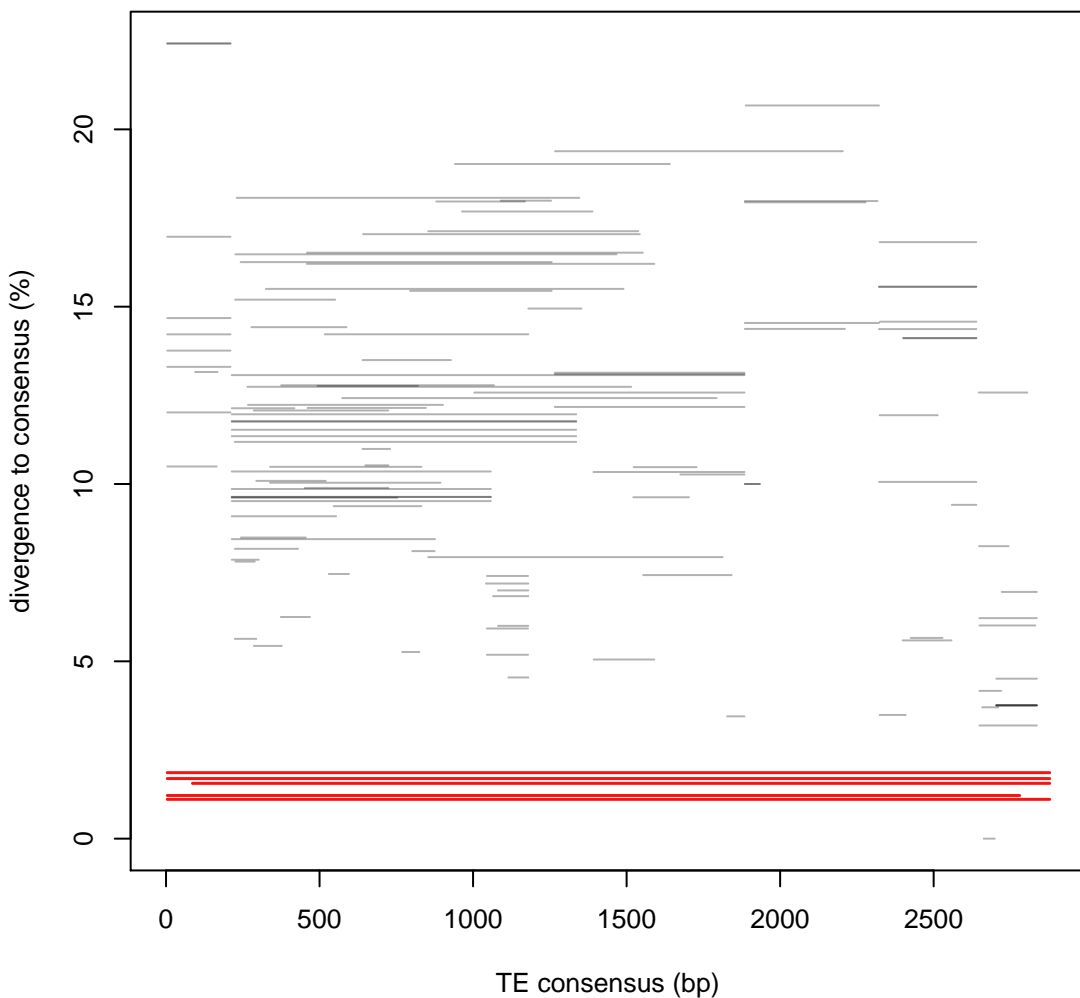
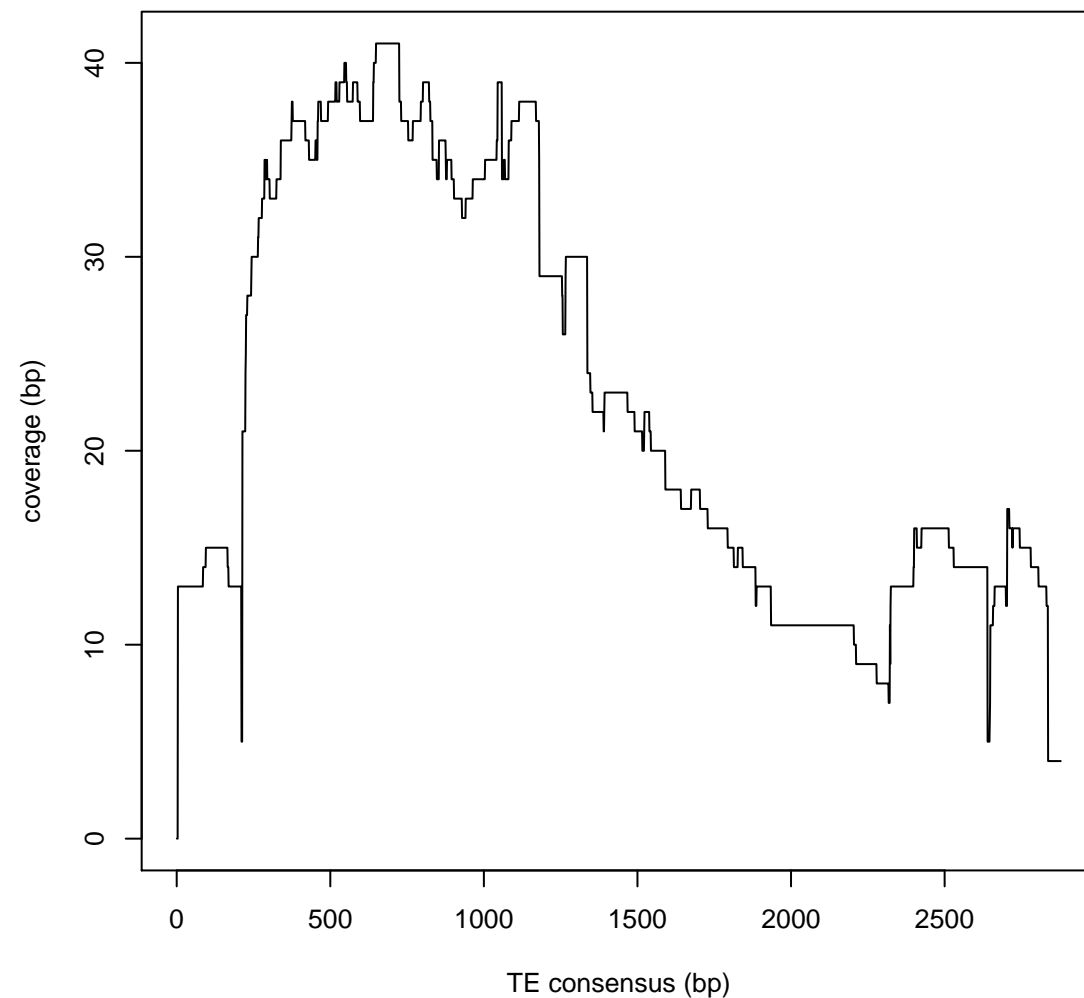


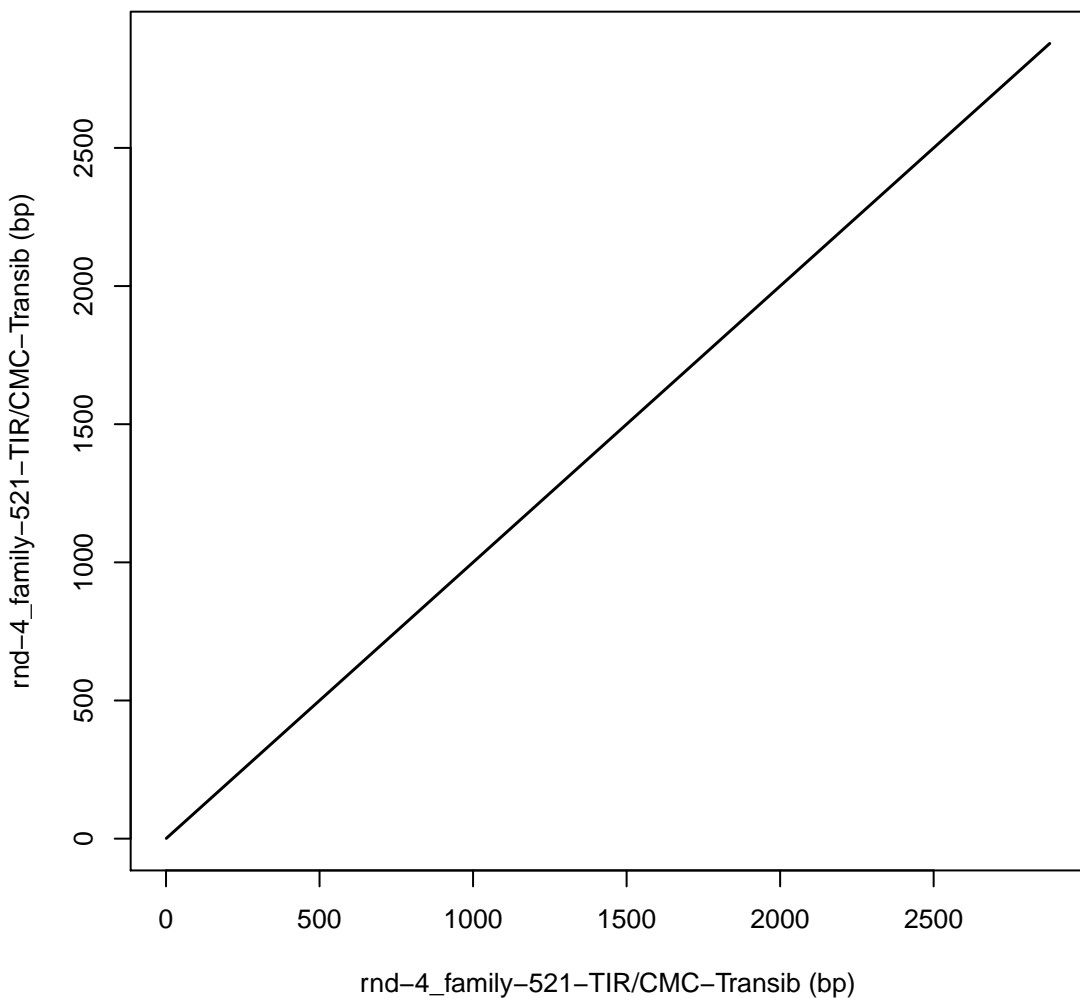
TE: rnd-4\_family-521-TIR/CMC-Transib  
consensus size: 2878bp; fragments: 128; full length: 5 ( $\geq 2590.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

