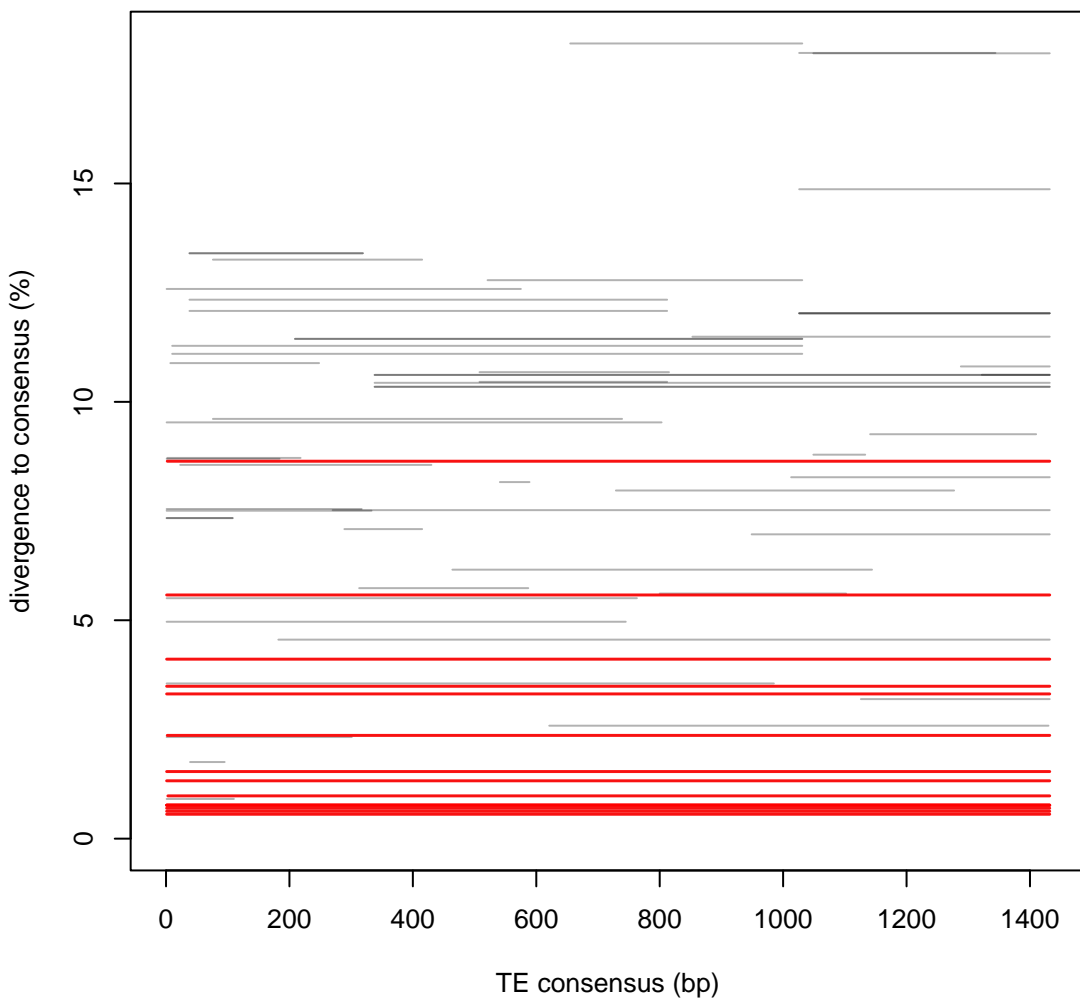
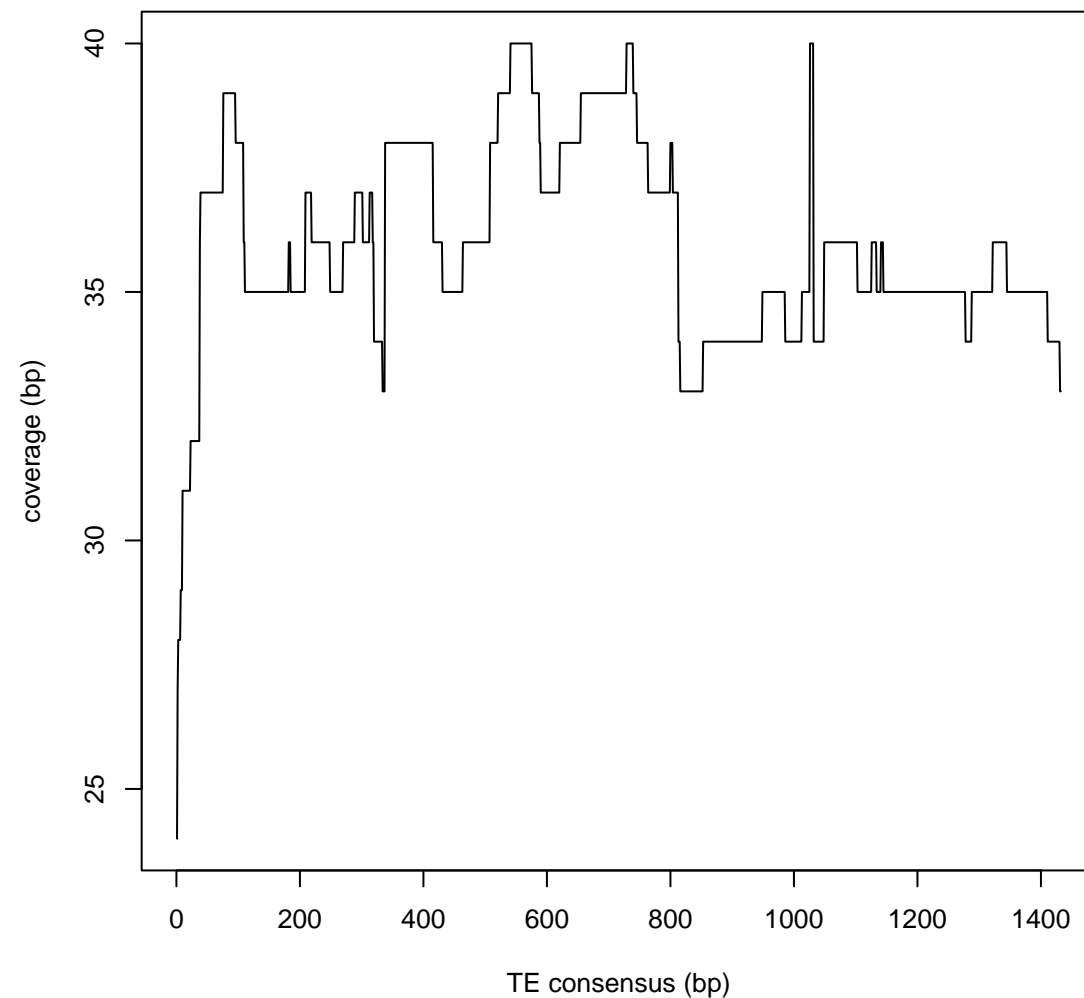


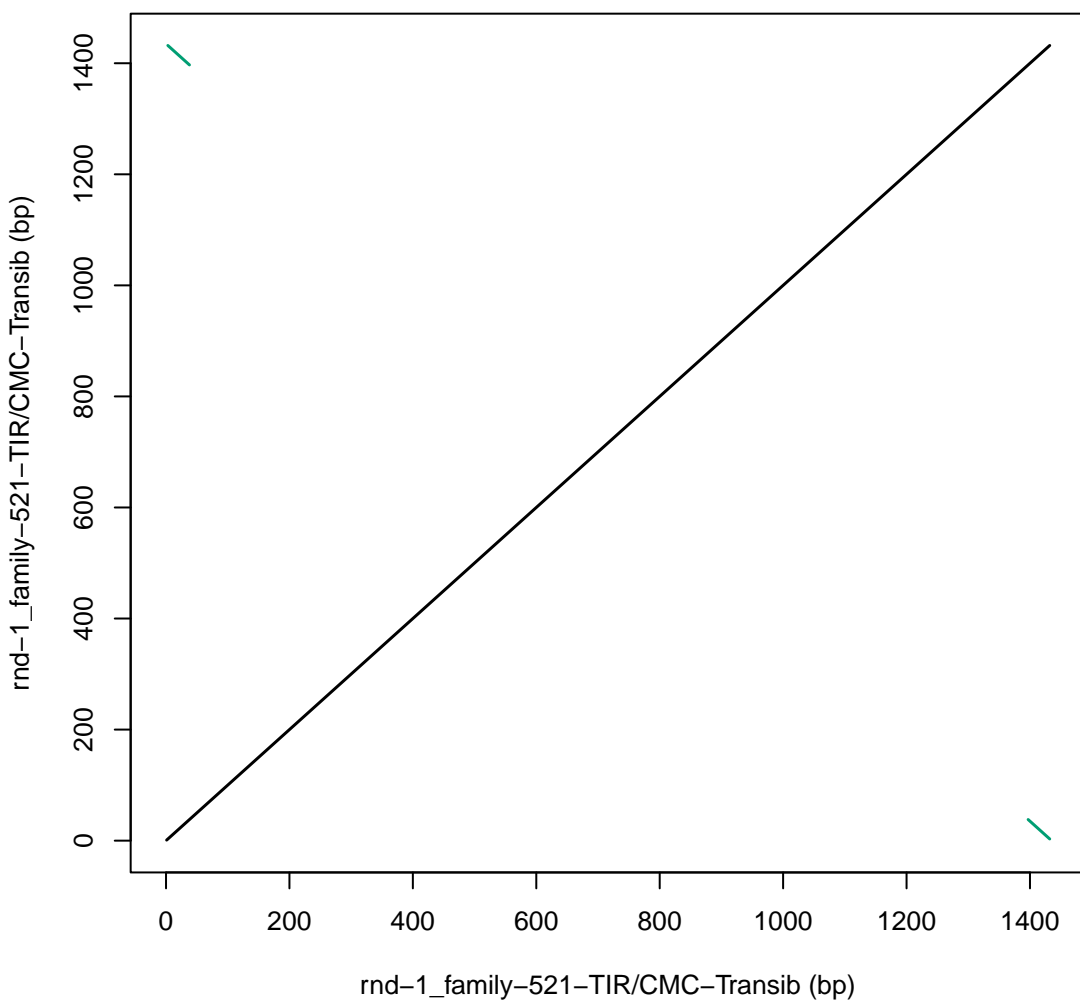
TE: rnd-1_family-521-TIR/CMC-Transib
consensus size: 1432bp; fragments: 73; full length: 15 ($\geq 1288.8\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

