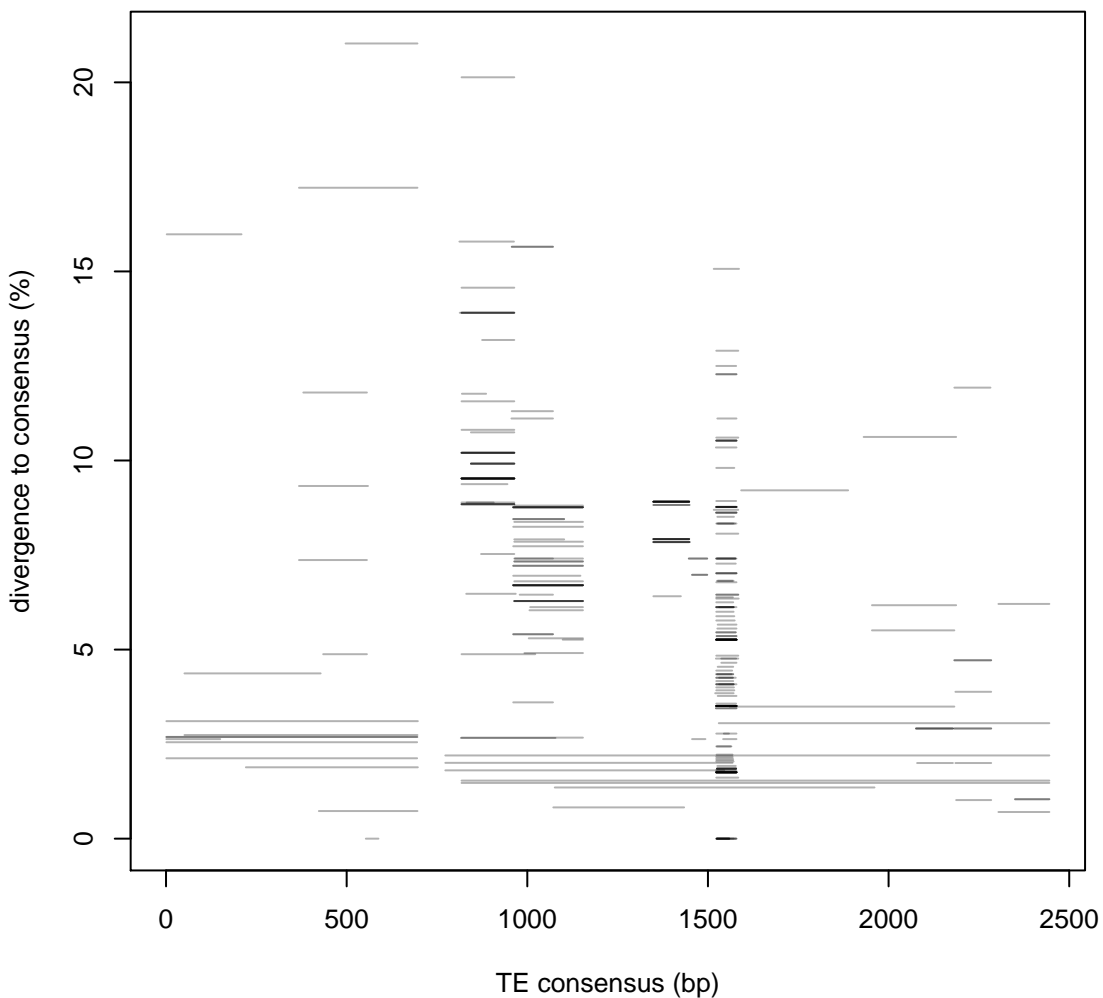
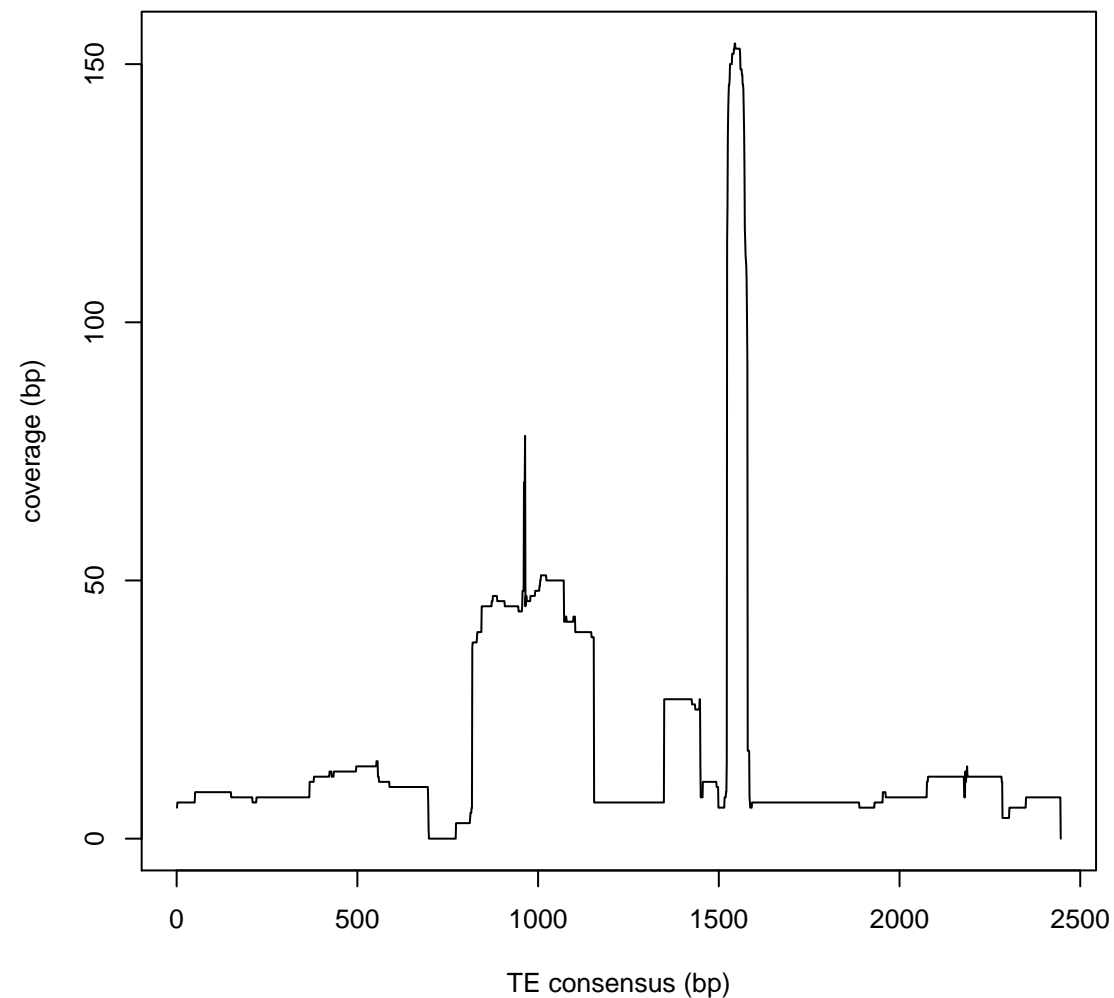


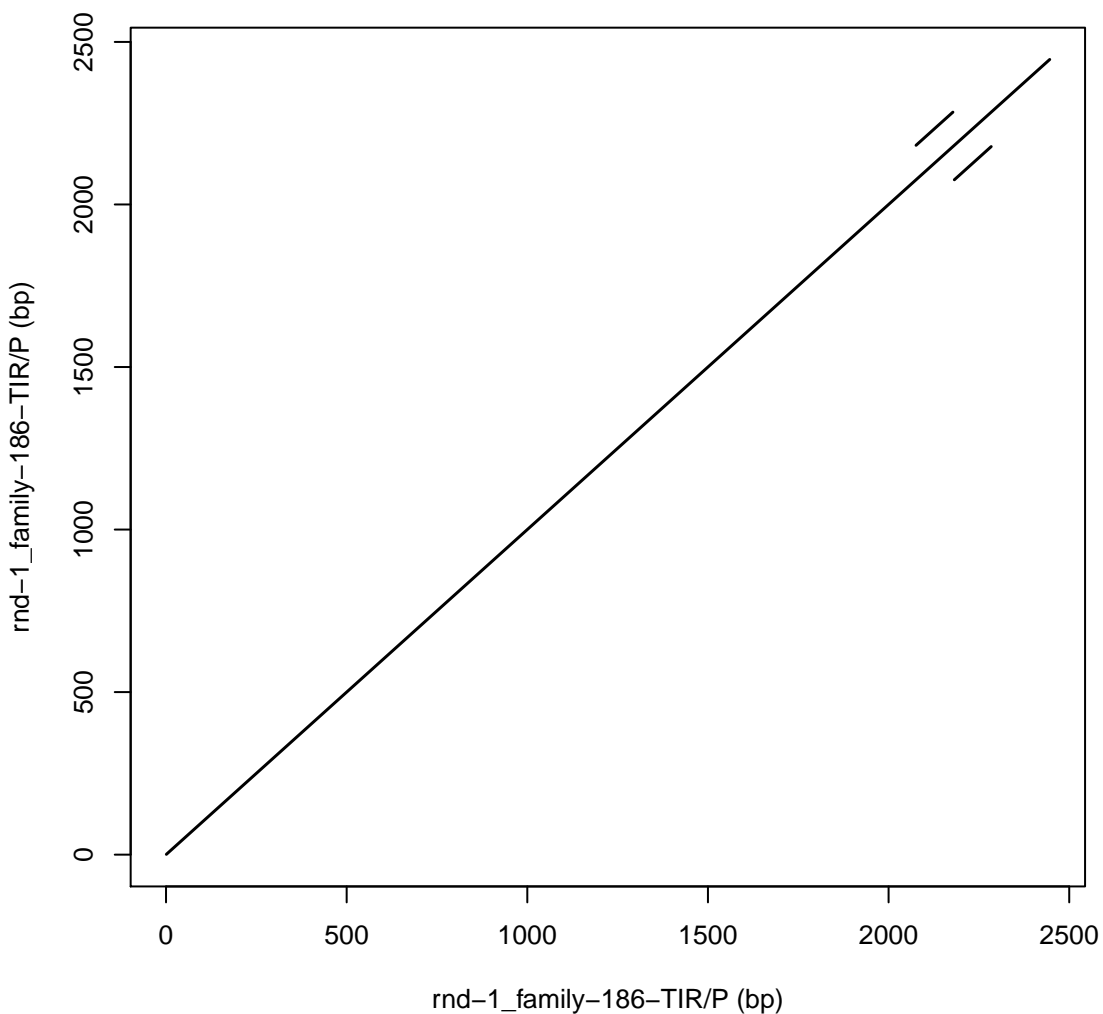
TE: rnd-1\_family-186-TIR/P  
consensus size: 2446bp; fragments: 305; full length: 0 (>=2201.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

