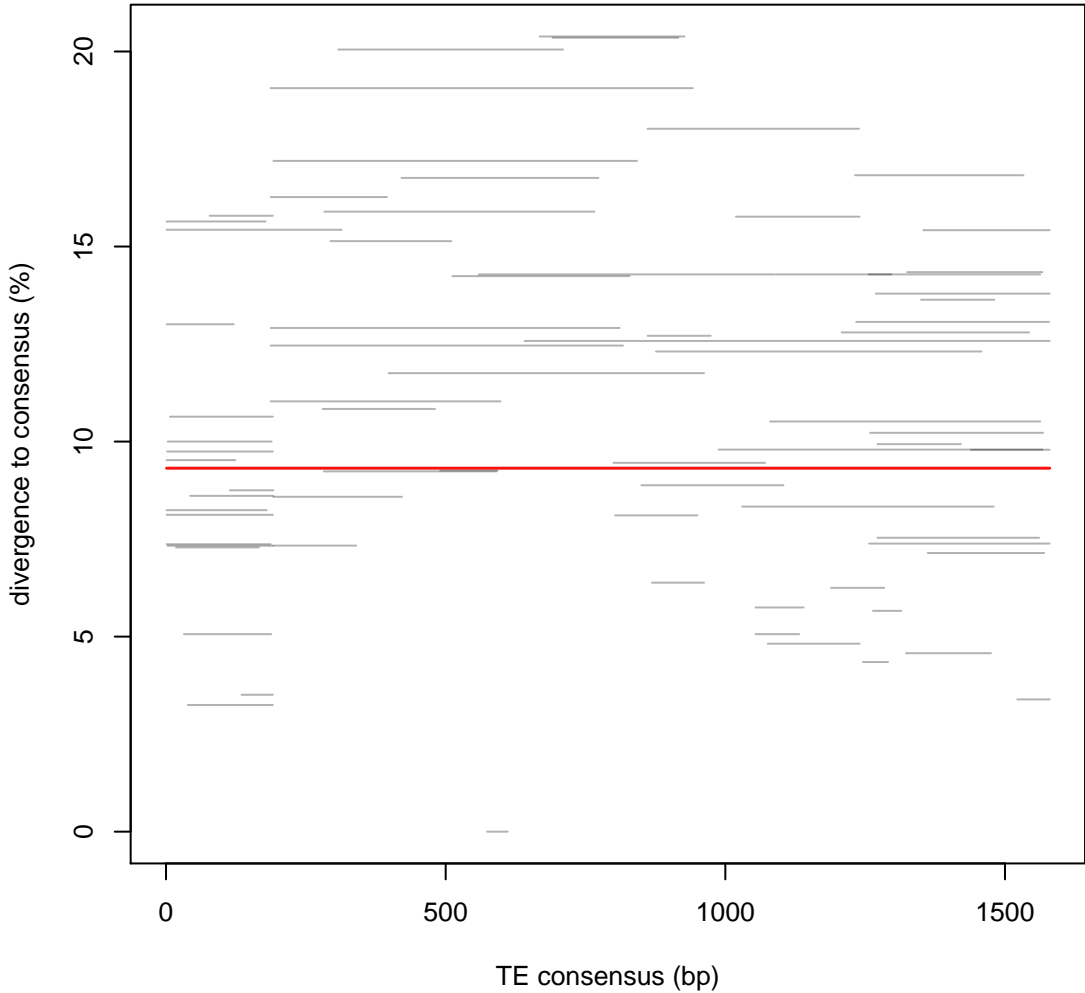
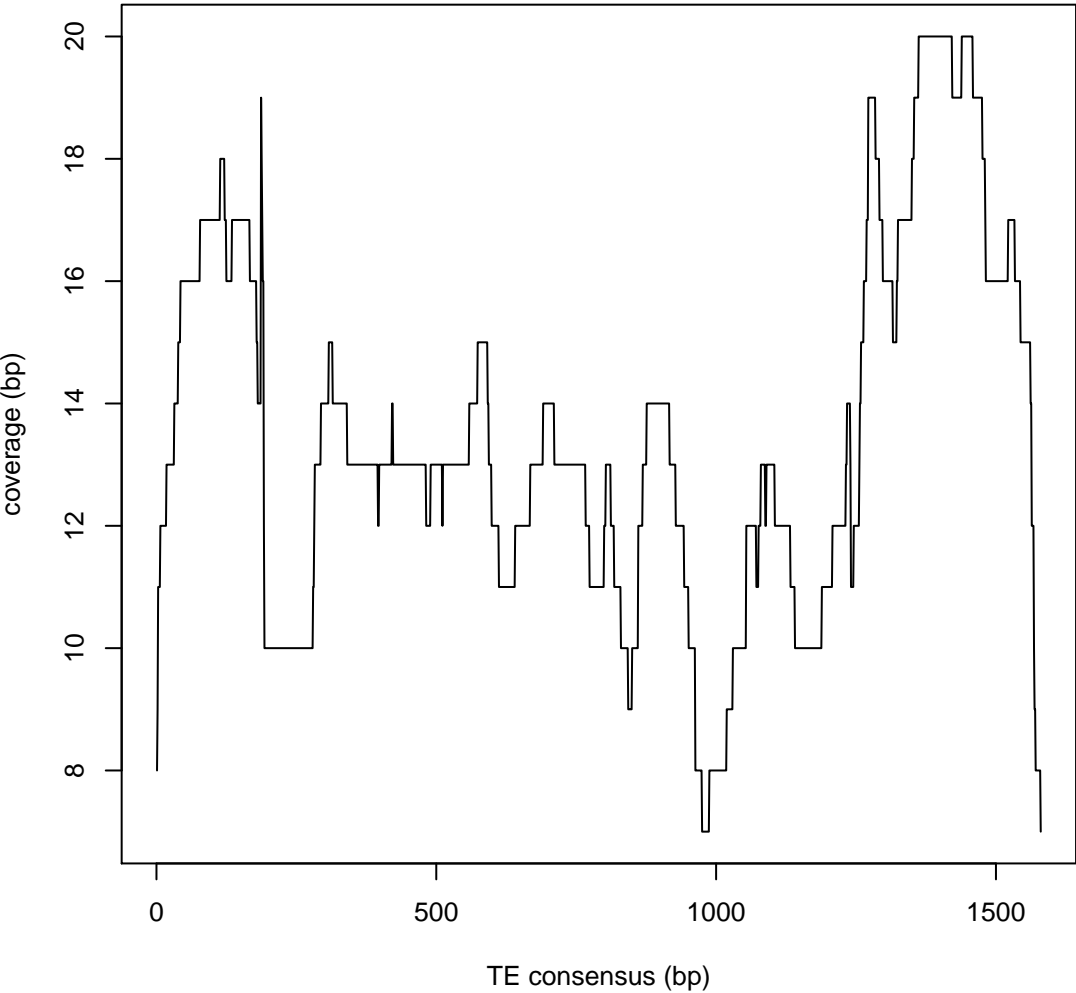


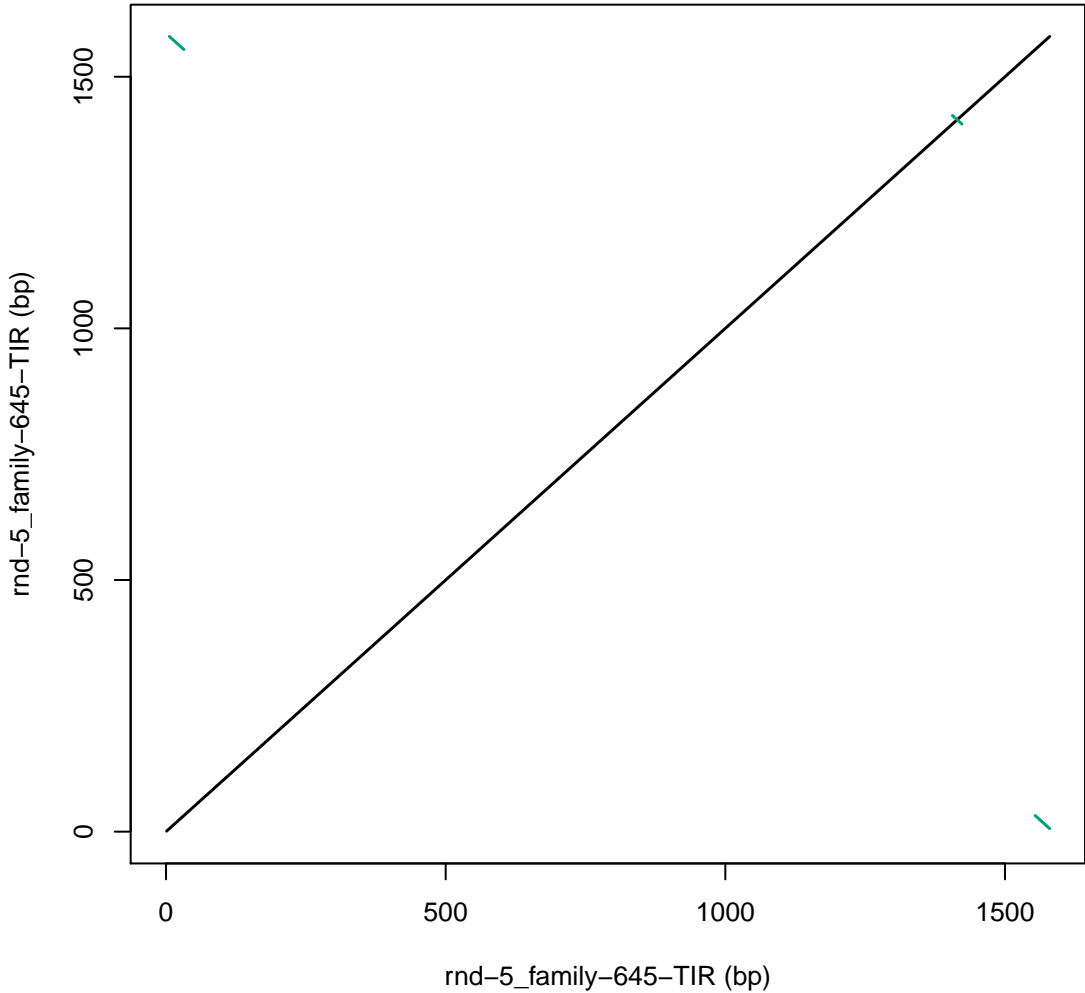
TE: rnd-5\_family-645-TIR  
consensus size: 1580bp; fragments: 75; full length: 1 (>=1422bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

