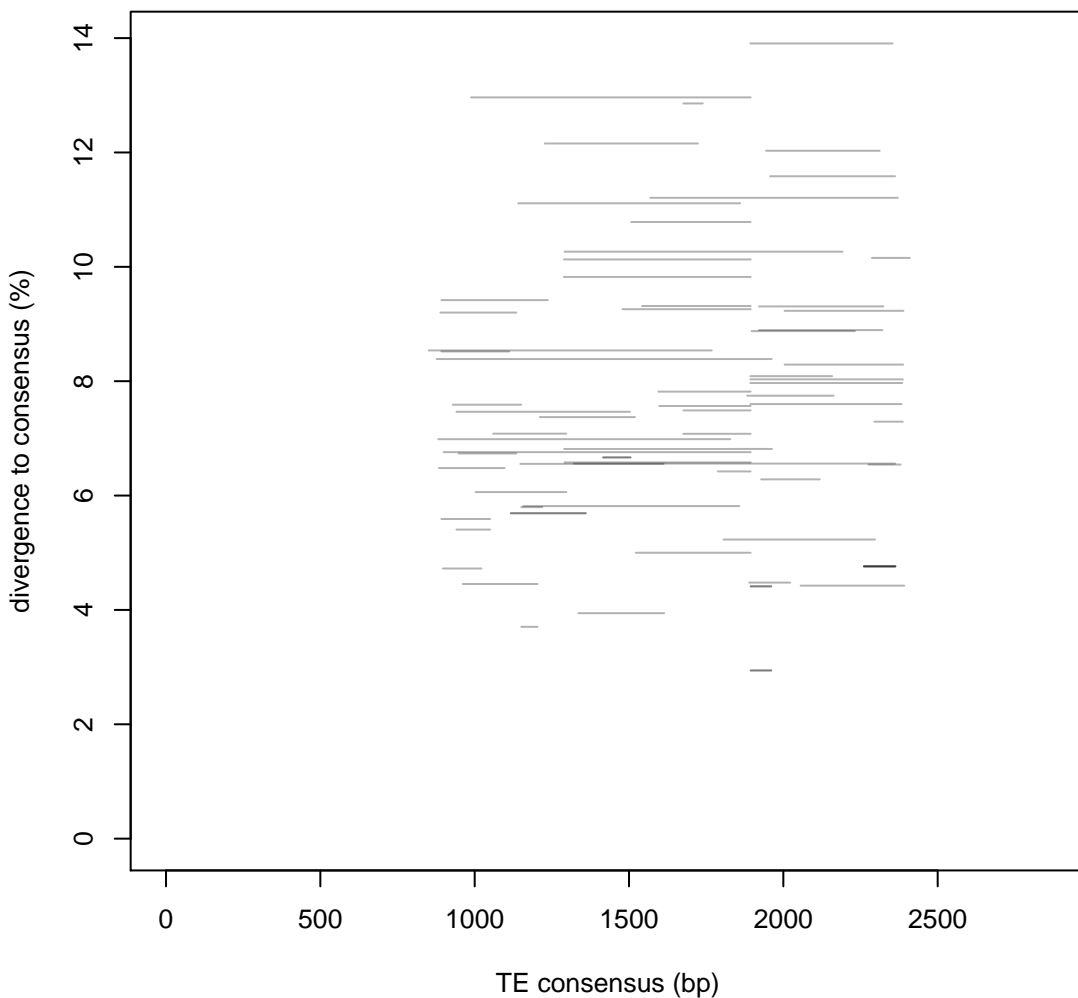
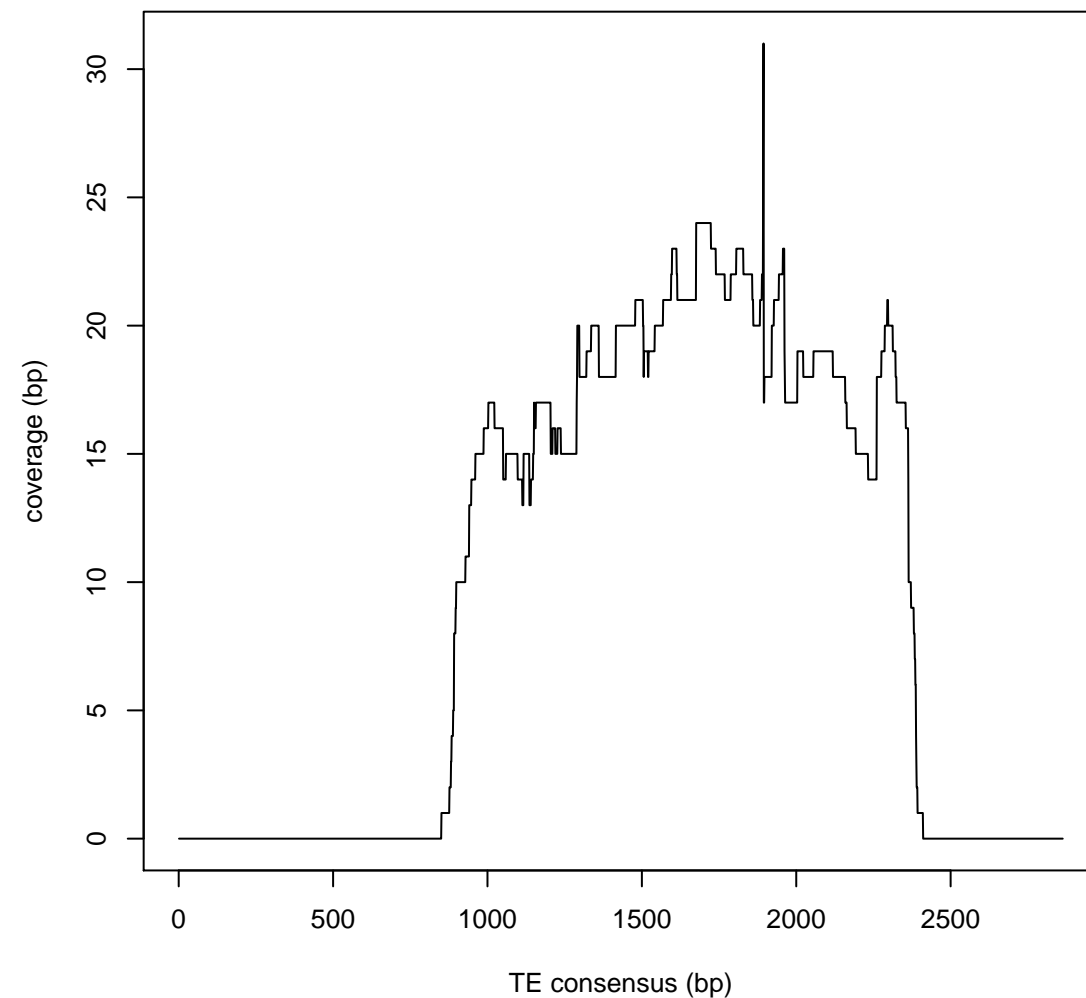


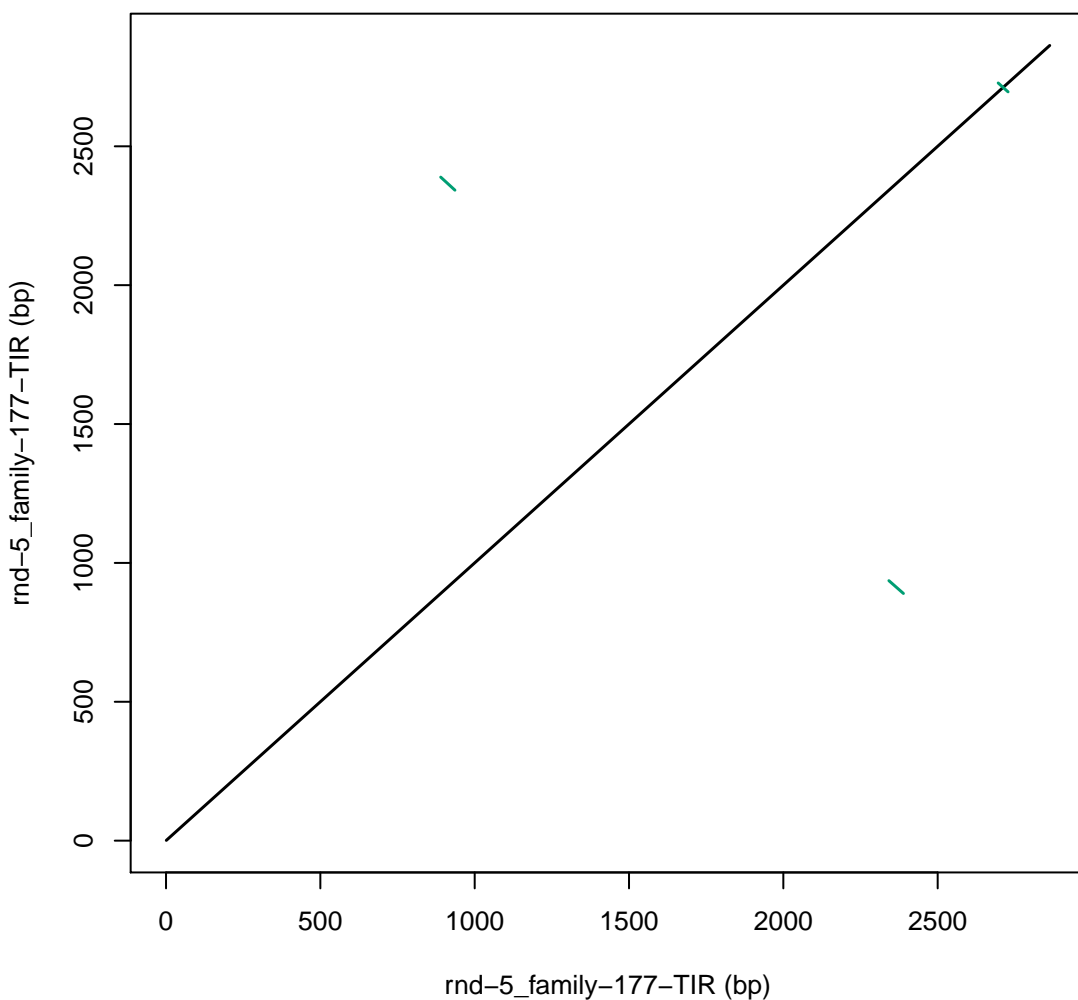
TE: rnd-5_family-177-TIR
consensus size: 2863bp; fragments: 75; full length: 0 (>=2576.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

