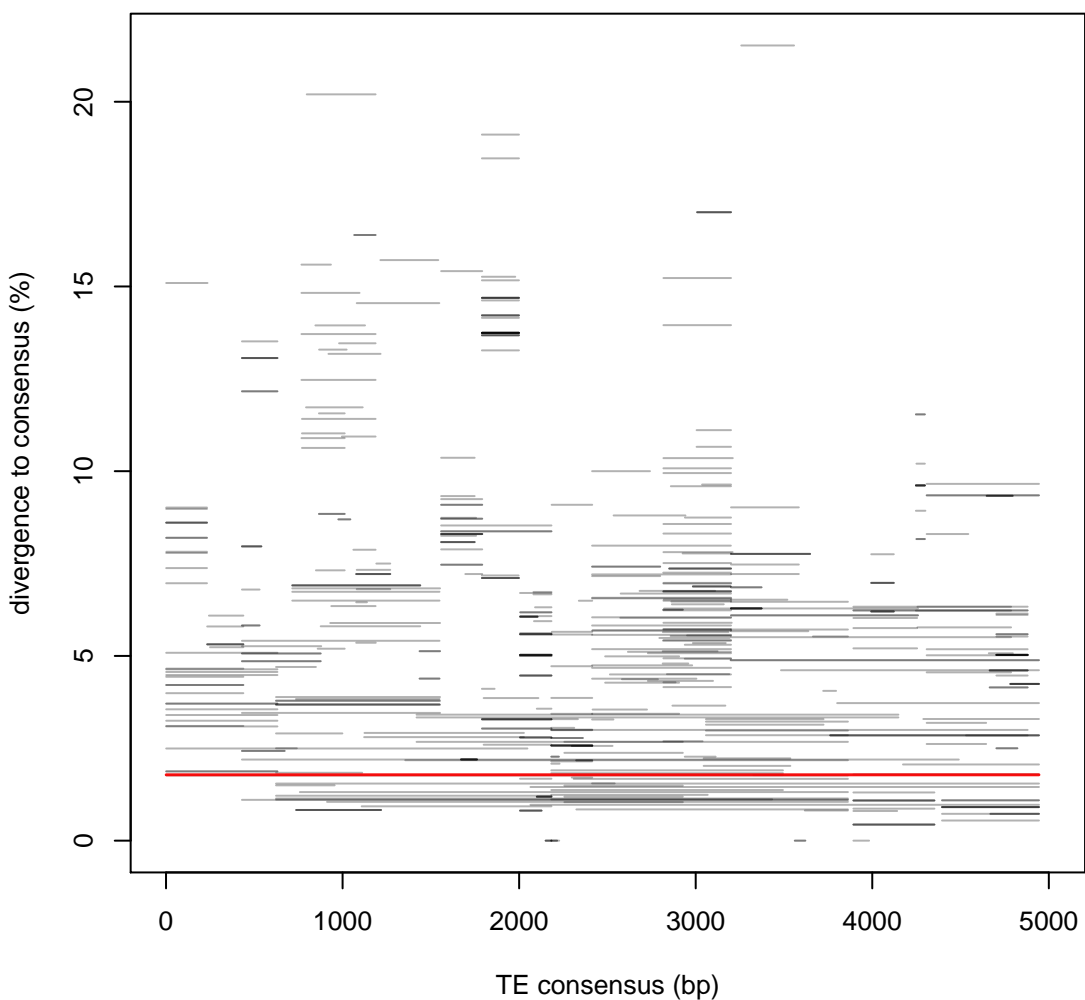
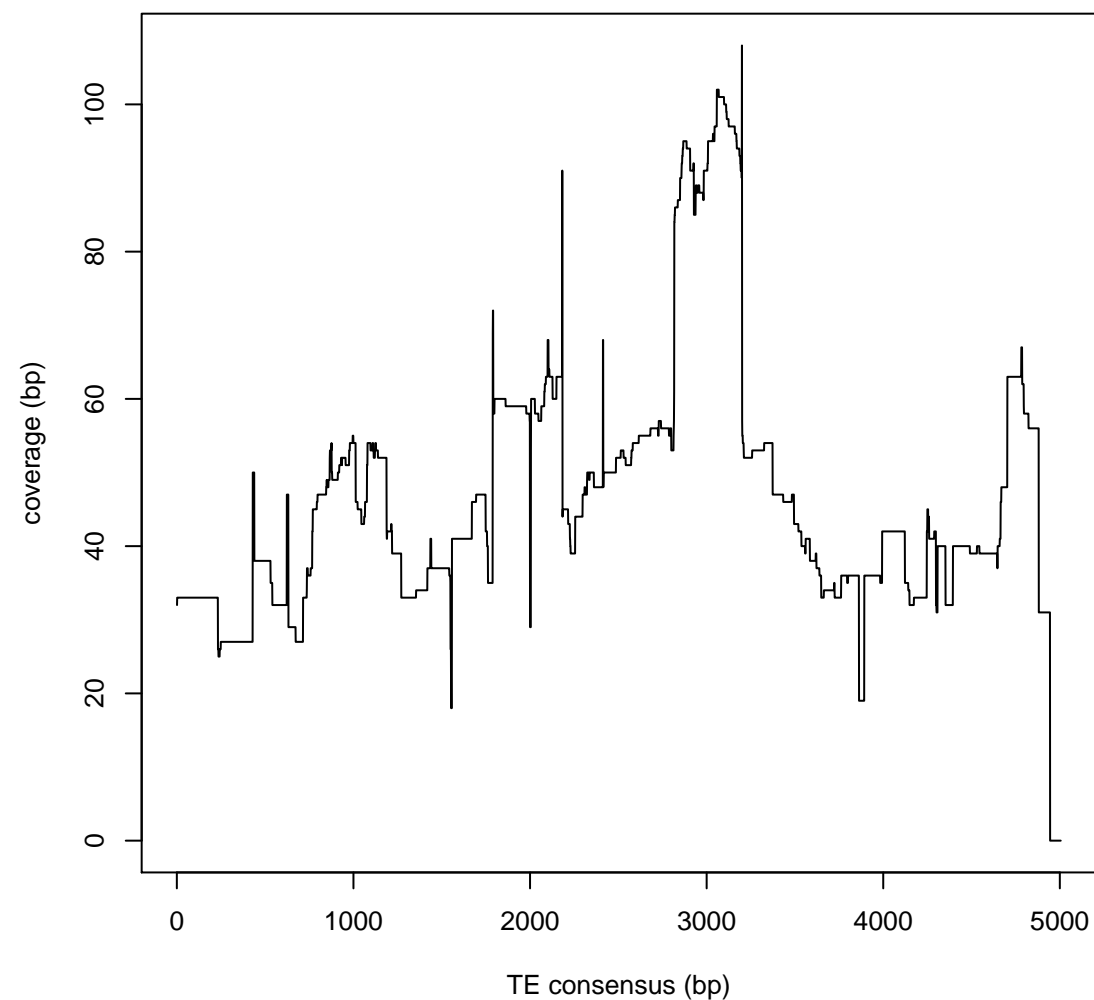


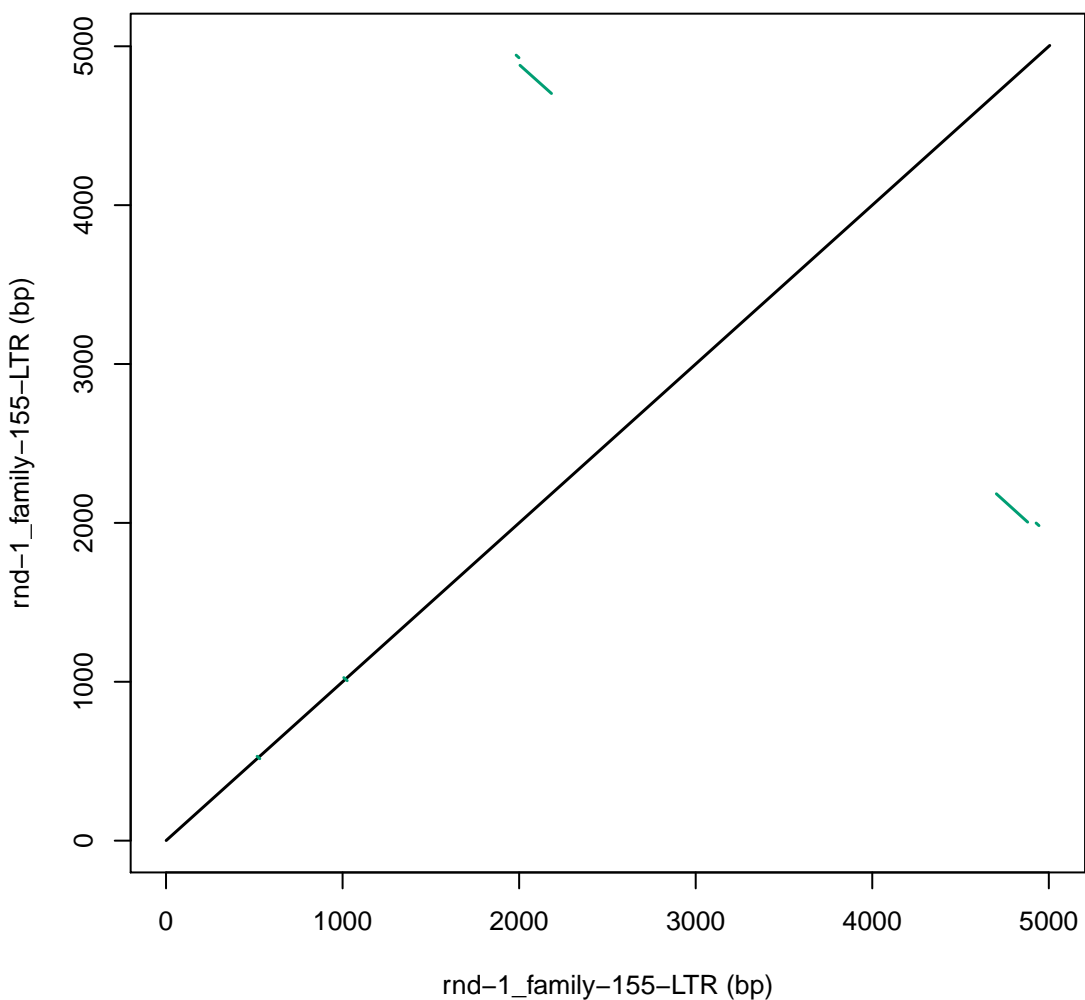
TE: rnd-1_family-155-LTR
consensus size: 5005bp; fragments: 516; full length: 1 (>=4504.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

