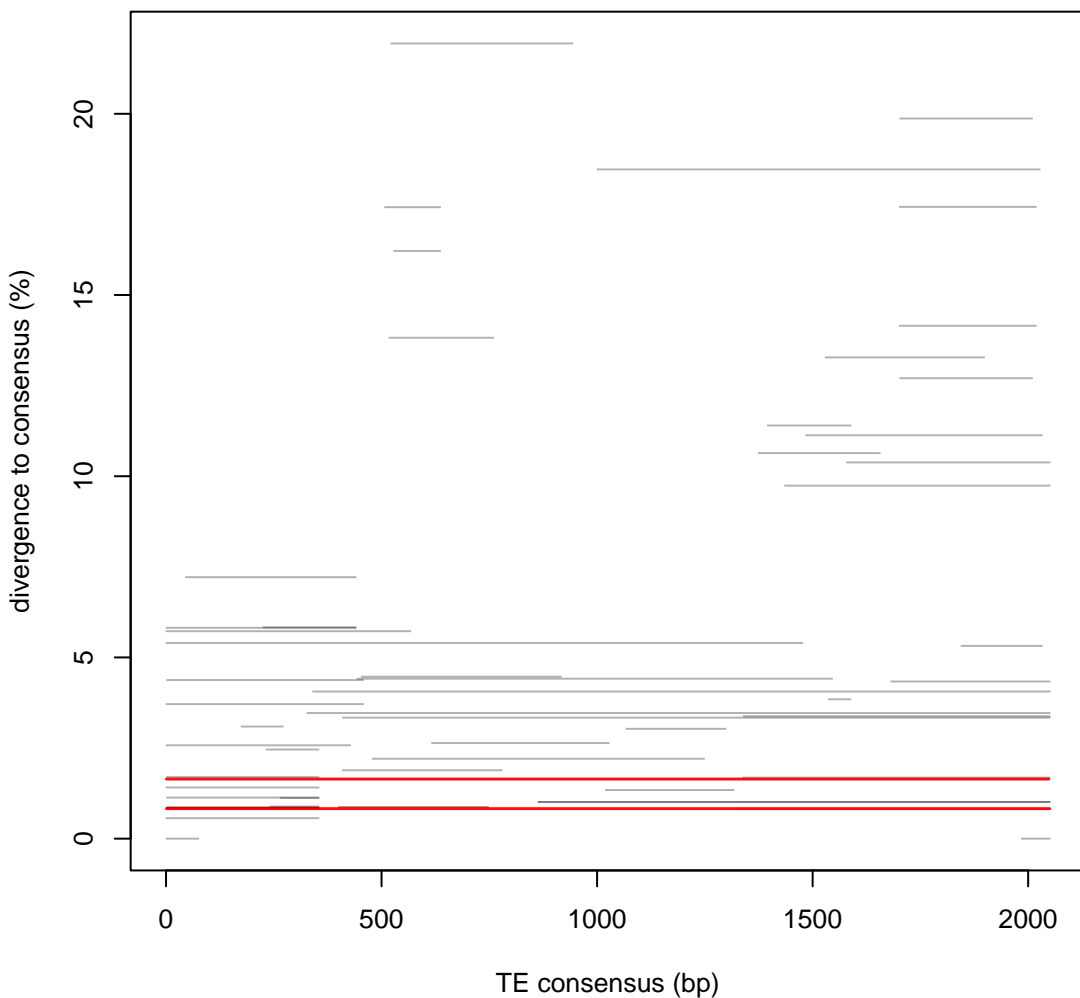
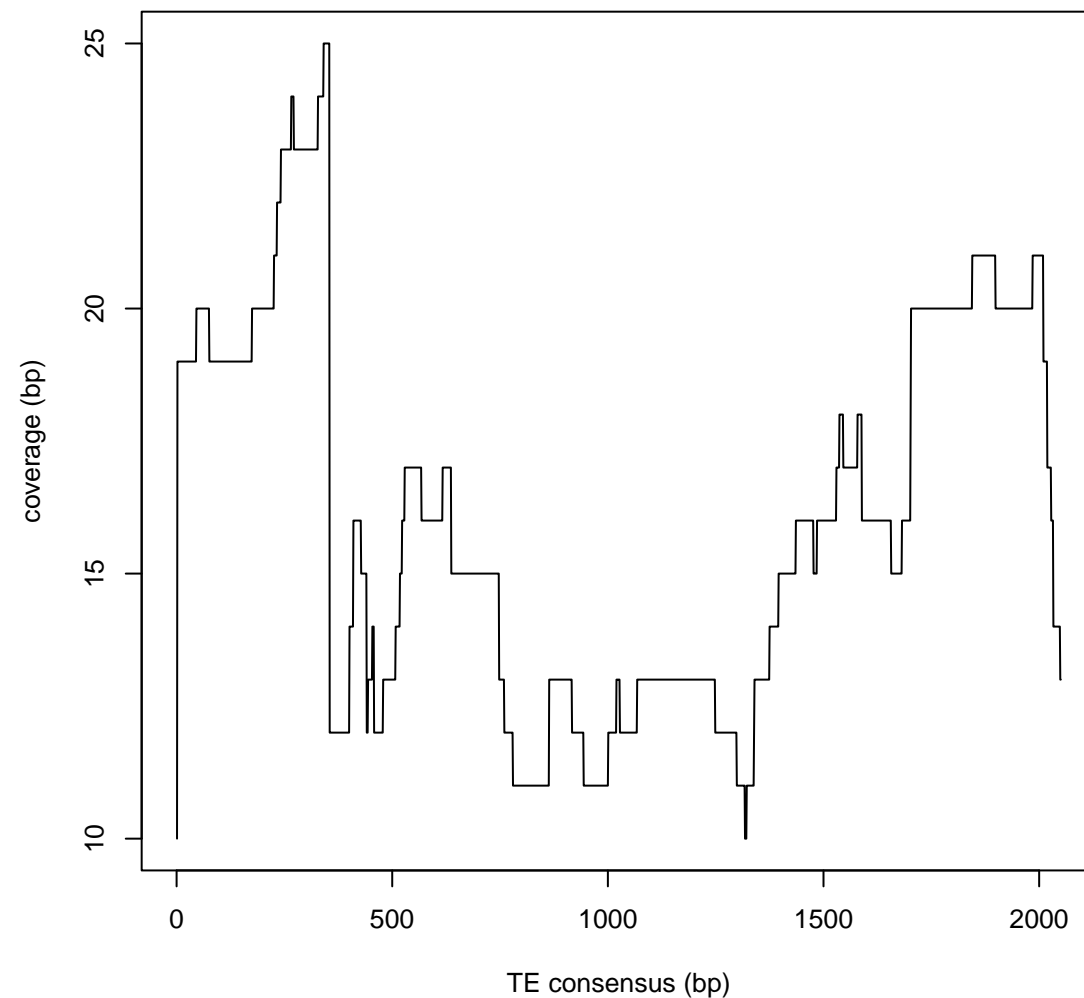


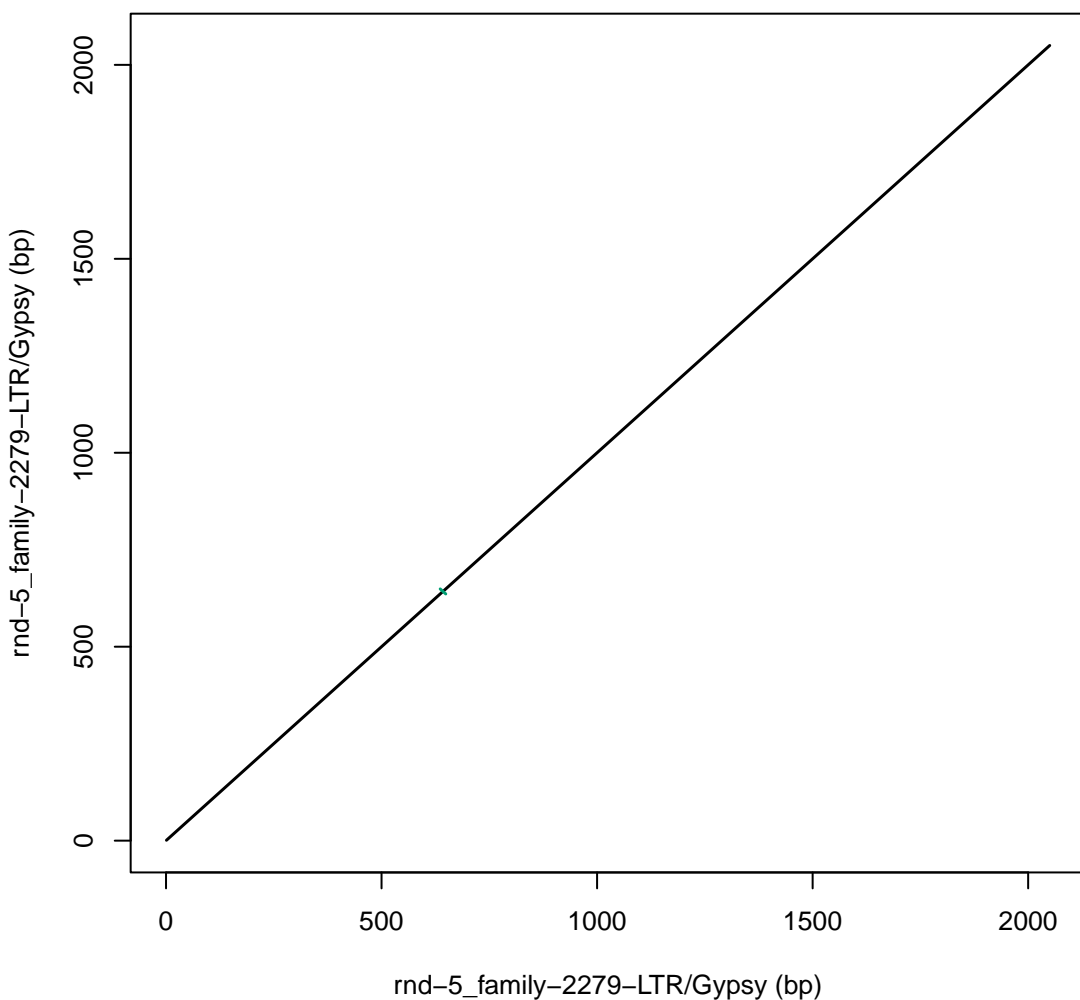
TE: rnd-5_family-2279-LTR/Gypsy
consensus size: 2050bp; fragments: 61; full length: 2 (>=1845bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

