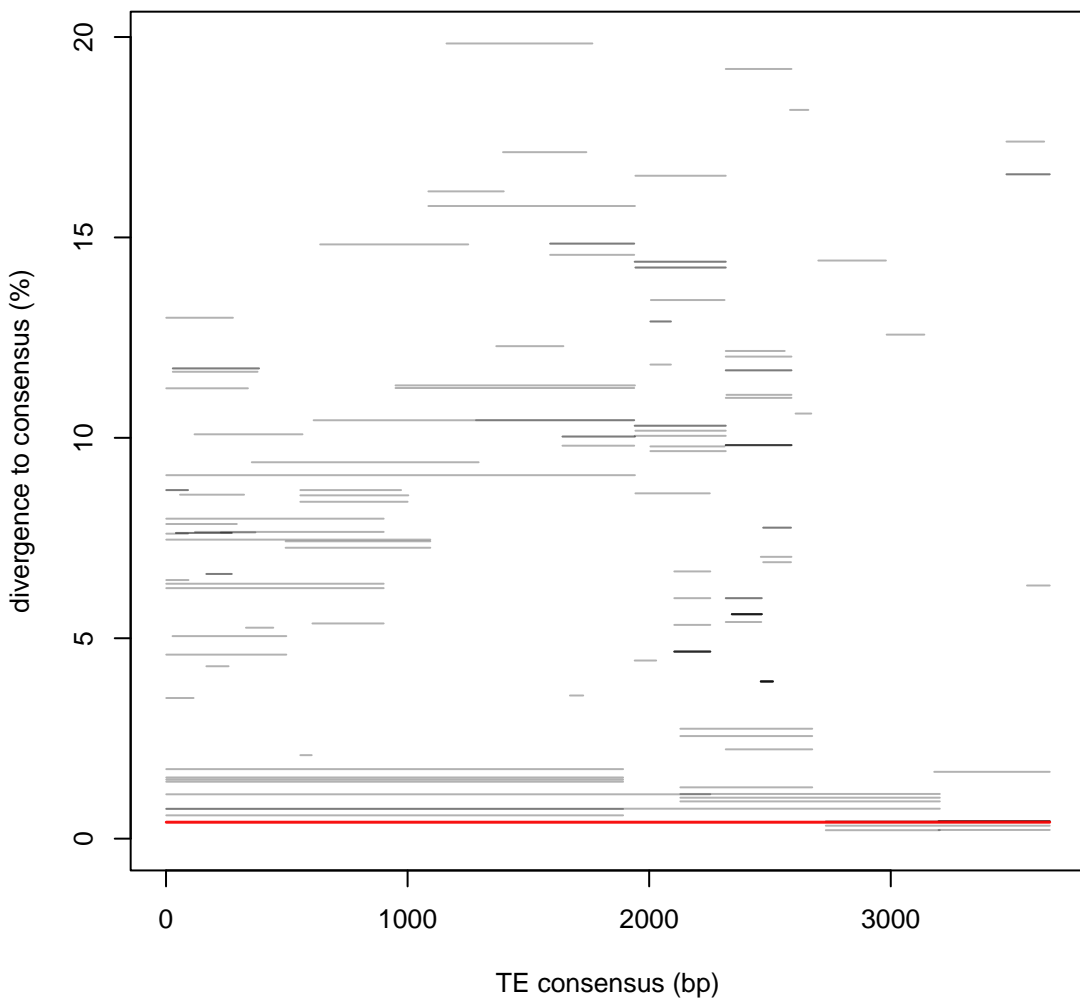
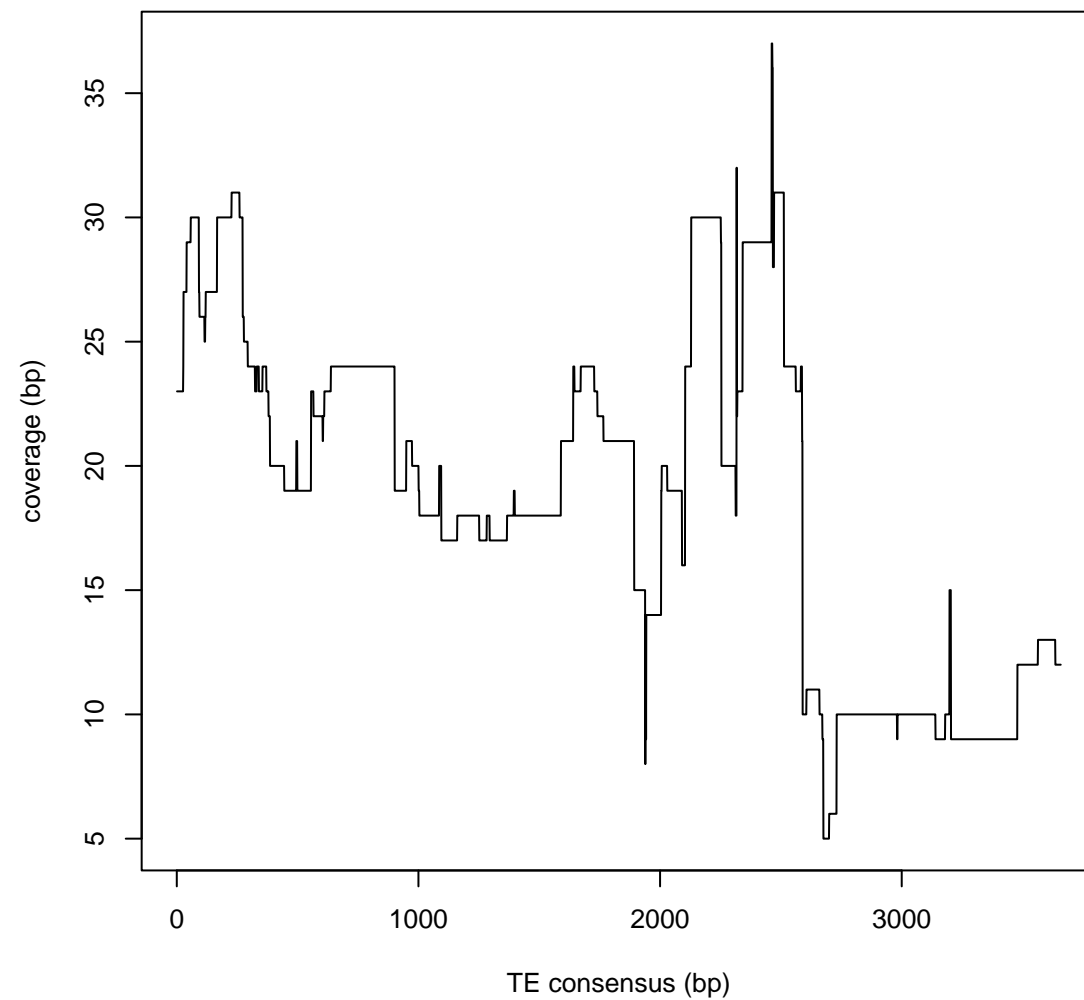


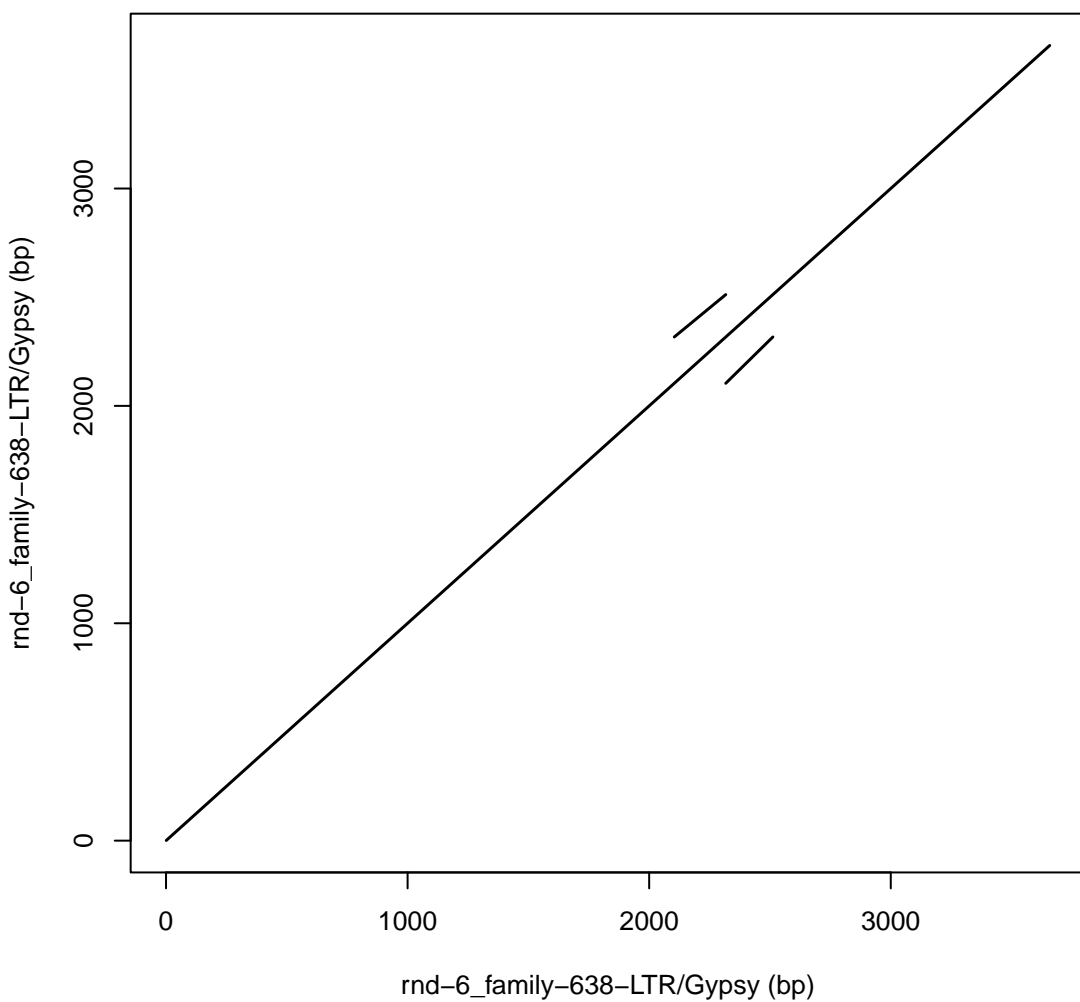
TE: rnd-6_family-638-LTR/Gypsy
consensus size: 3658bp; fragments: 143; full length: 1 (>=3292.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

