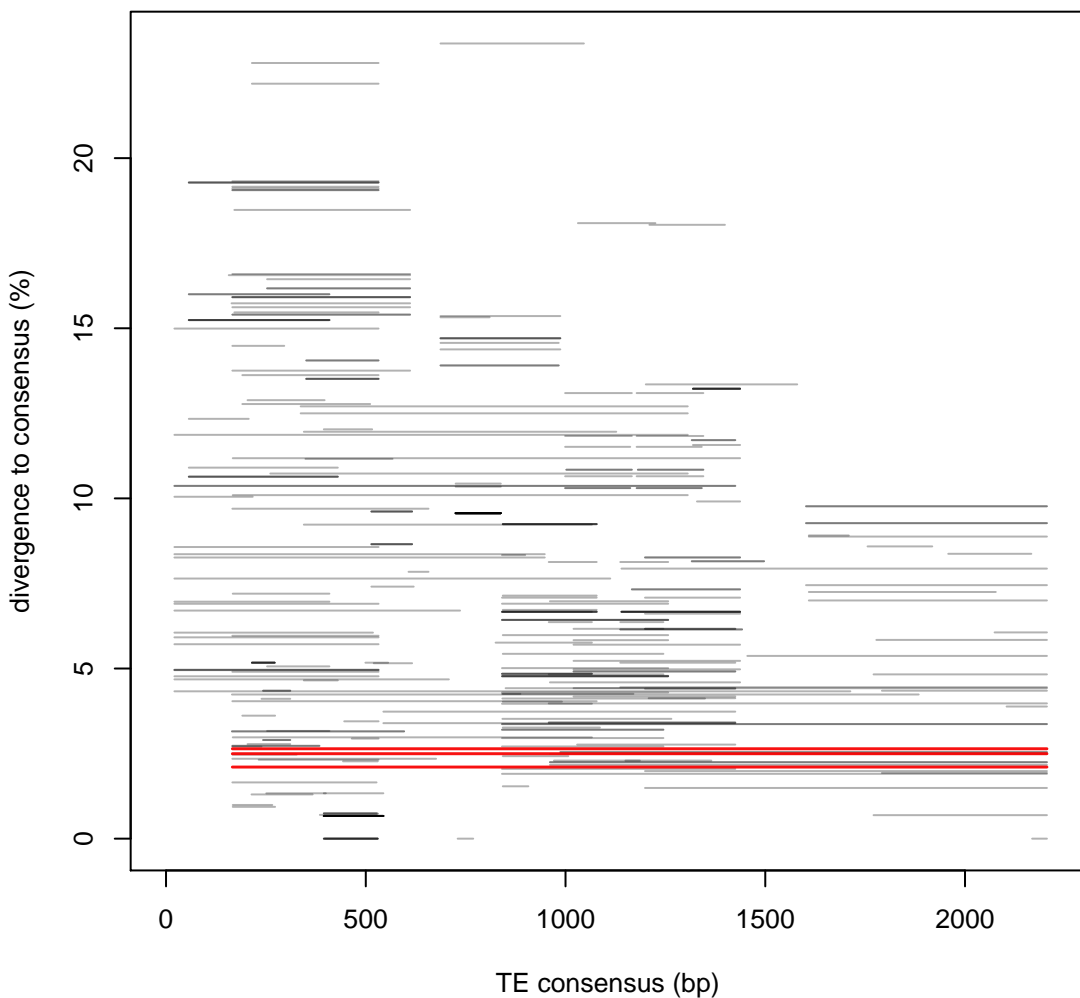
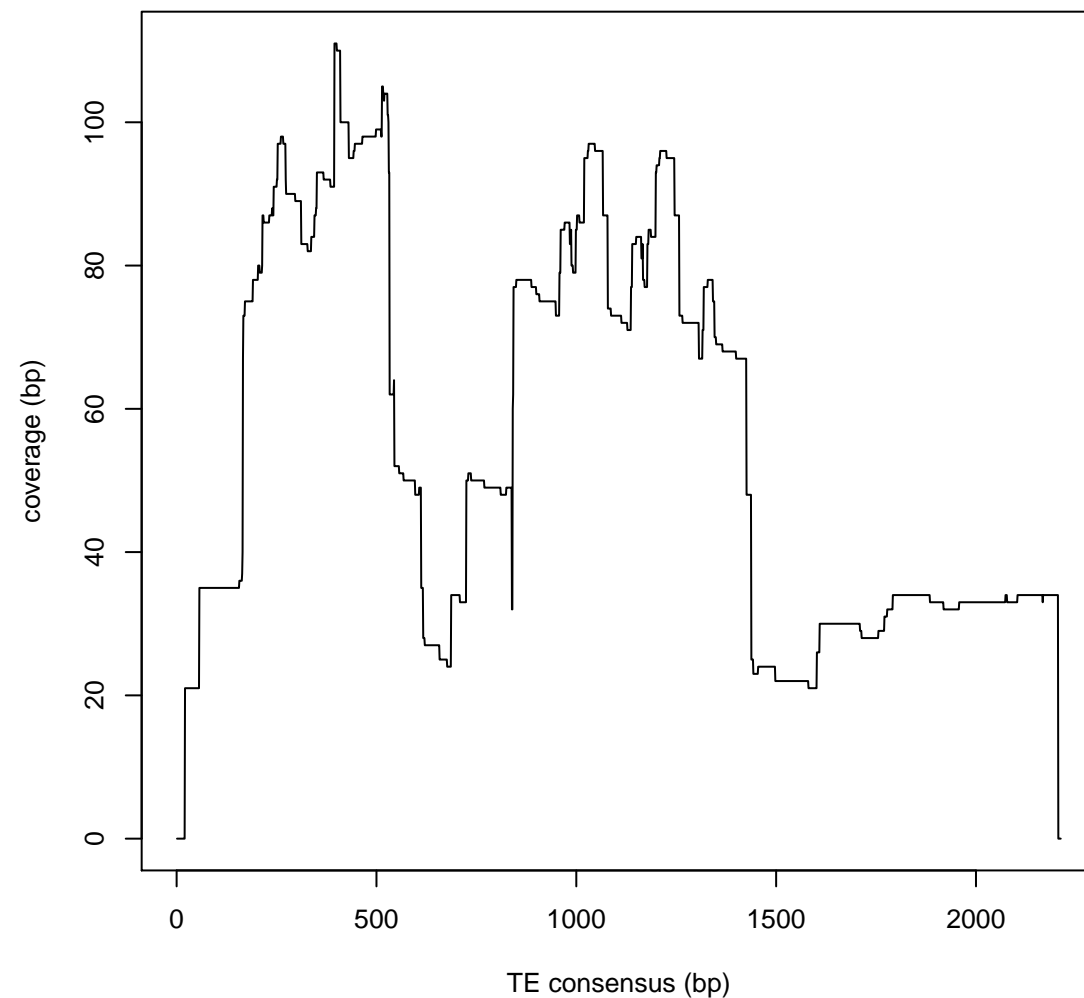


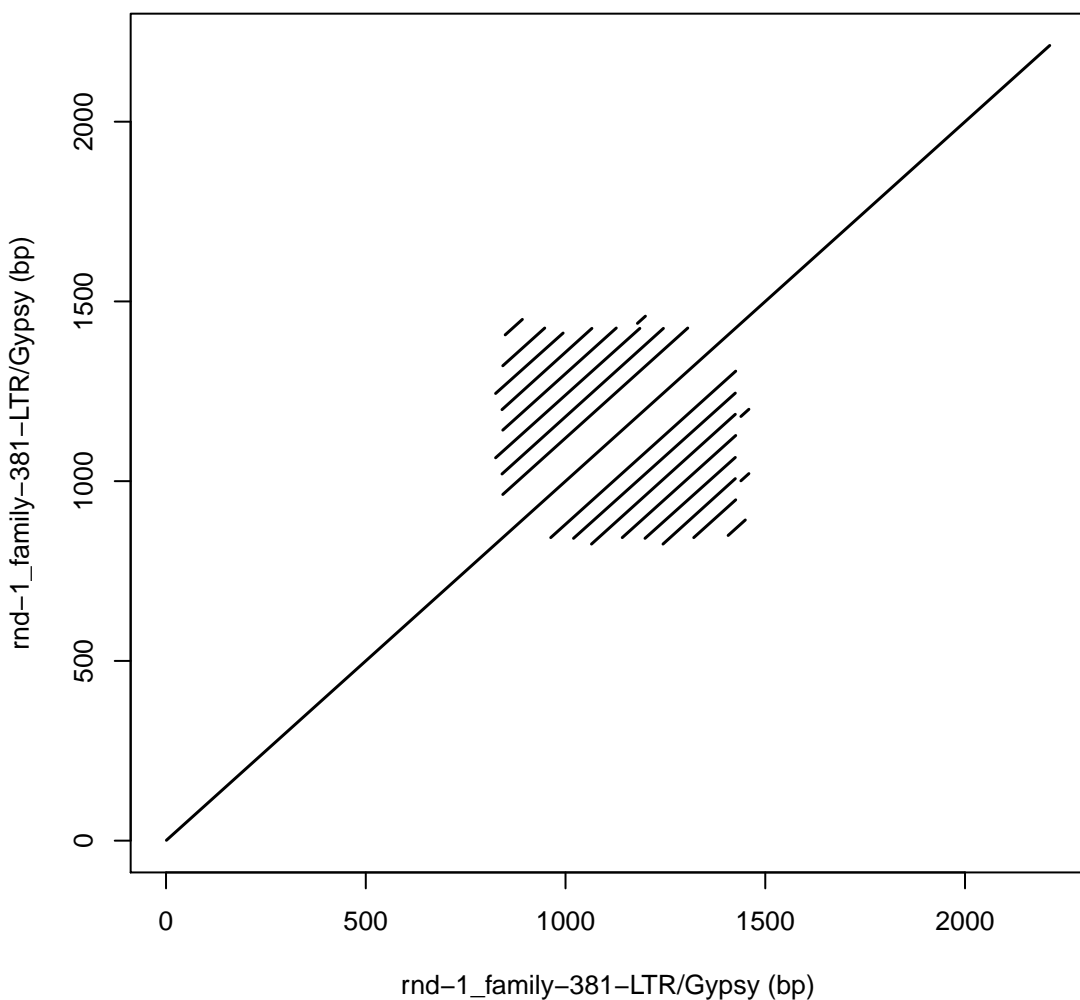
TE: rnd-1_family-381-LTR/Gypsy
consensus size: 2212bp; fragments: 324; full length: 3 (≥ 1990.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

