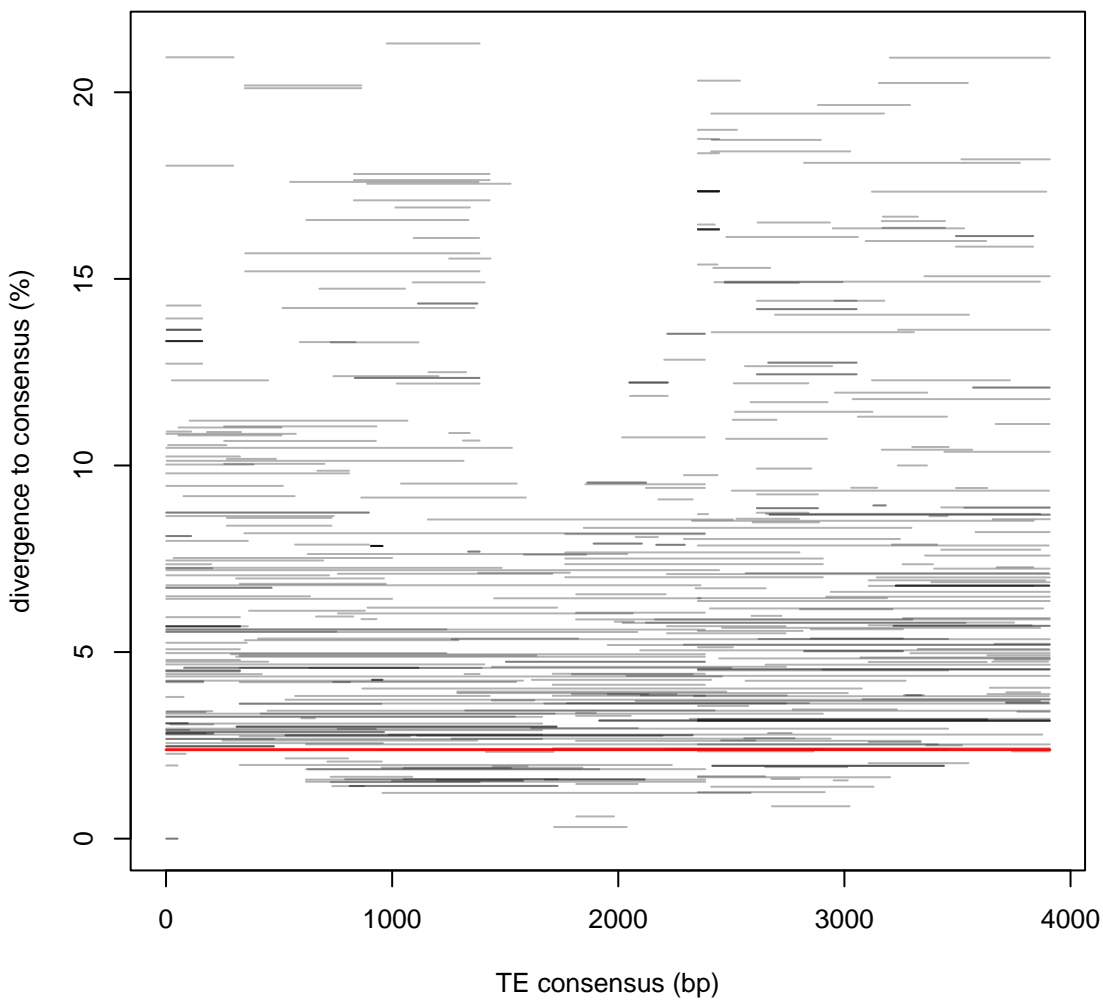
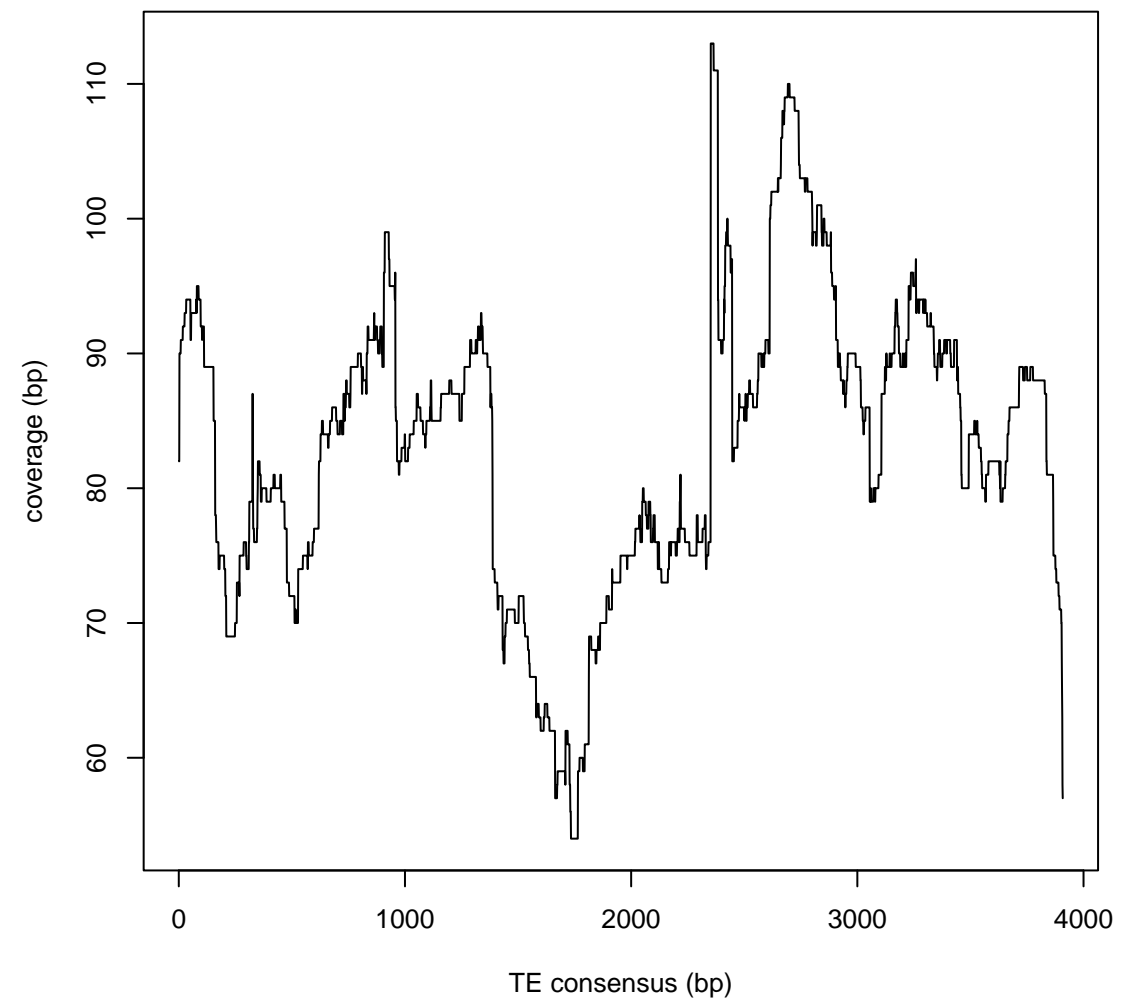


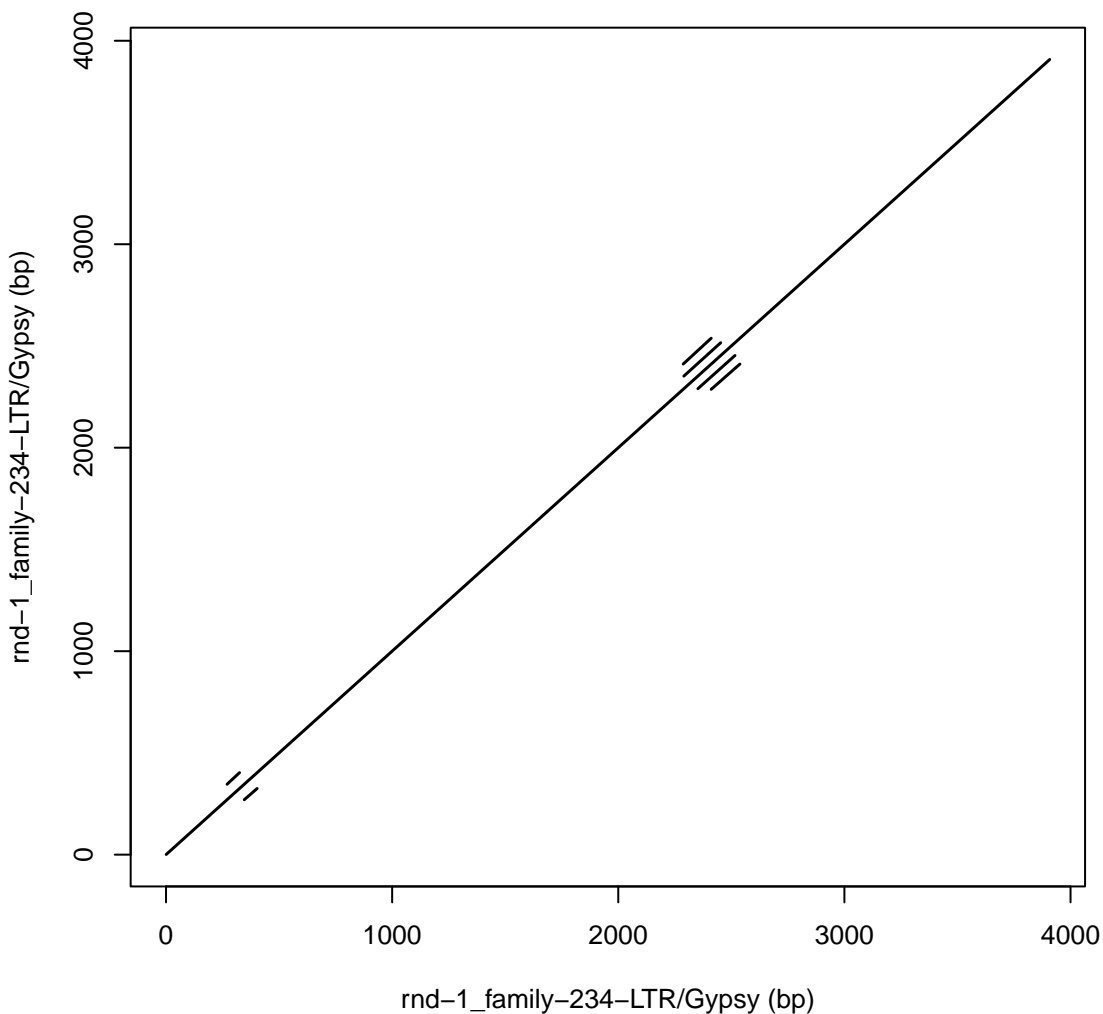
TE: rnd-1_family-234-LTR/Gypsy
consensus size: 3908bp; fragments: 469; full length: 1 (≥ 3517.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

