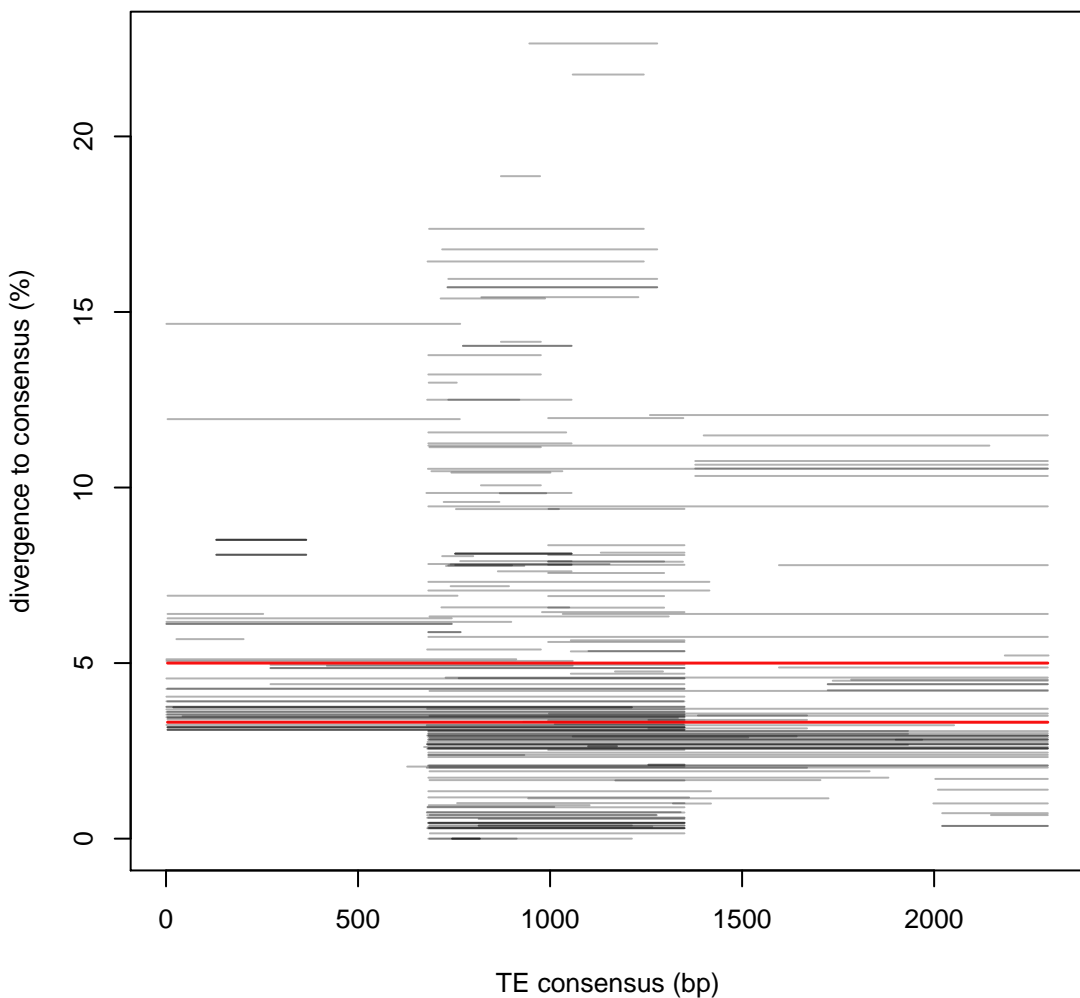
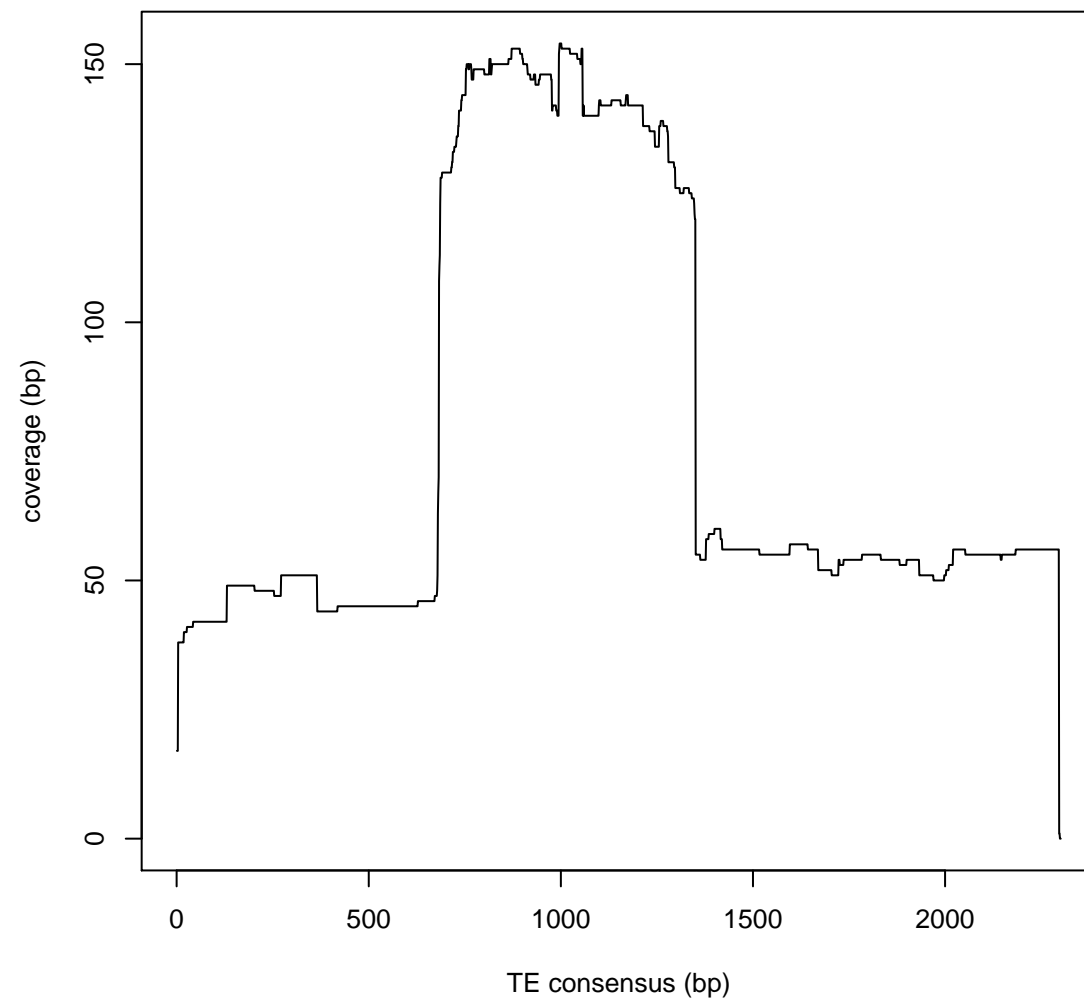


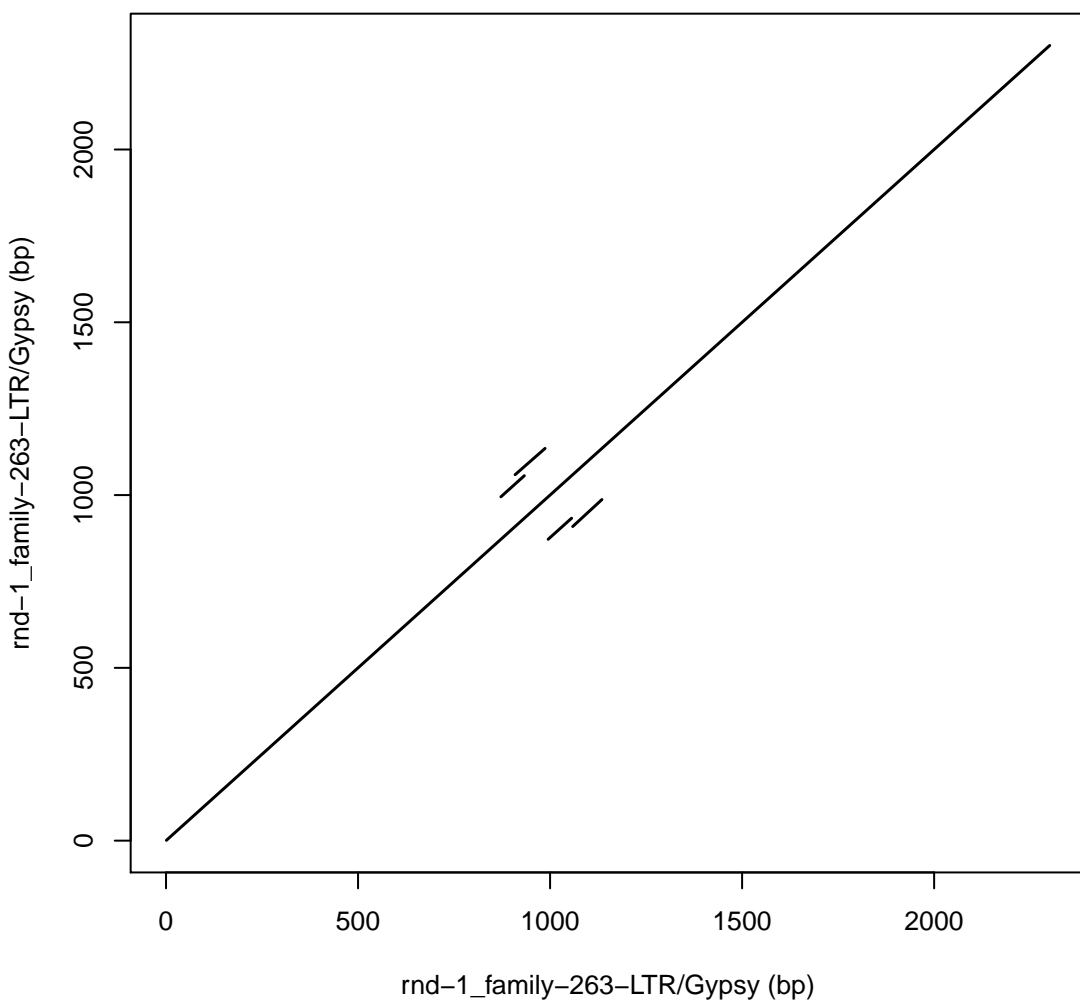
TE: rnd-1_family-263-LTR/Gypsy
consensus size: 2301bp; fragments: 235; full length: 2 (≥ 2070.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

