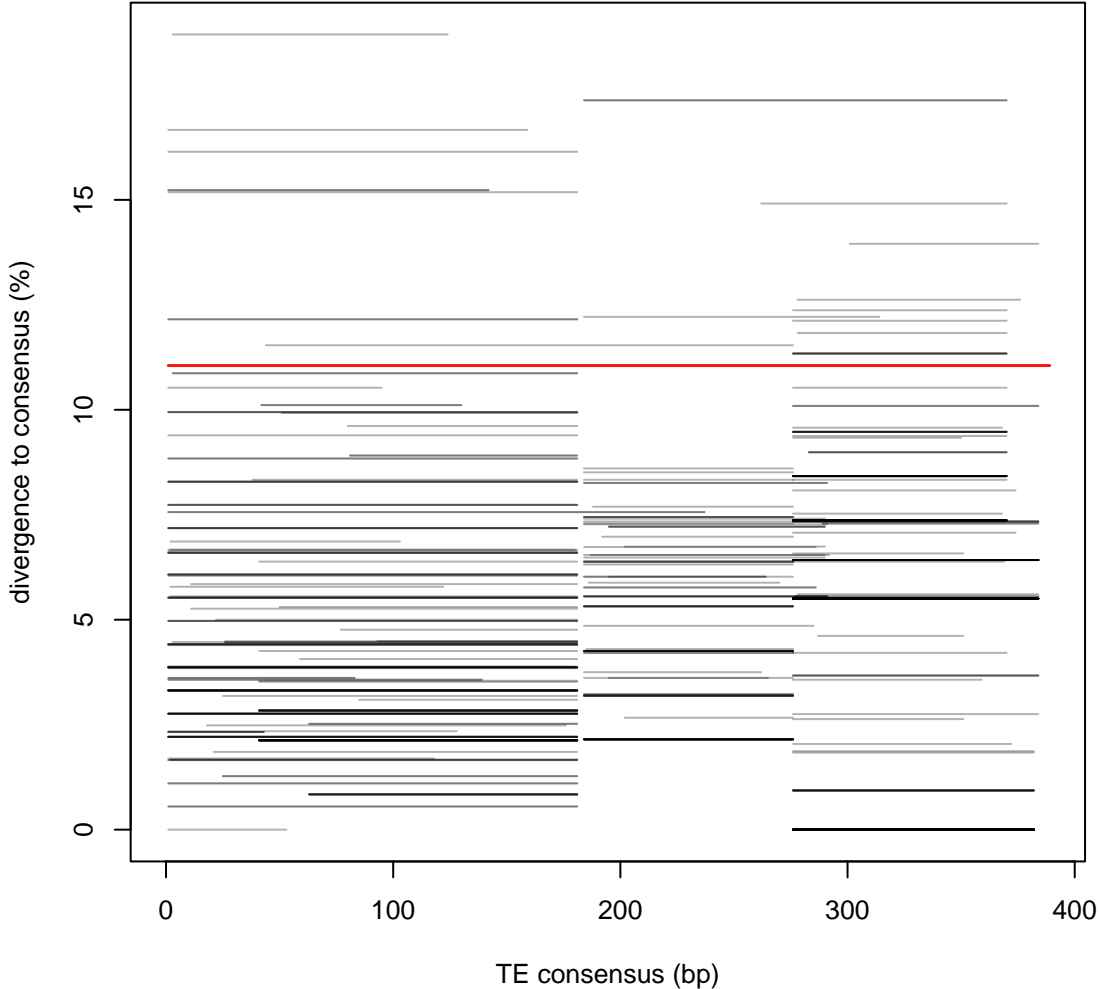
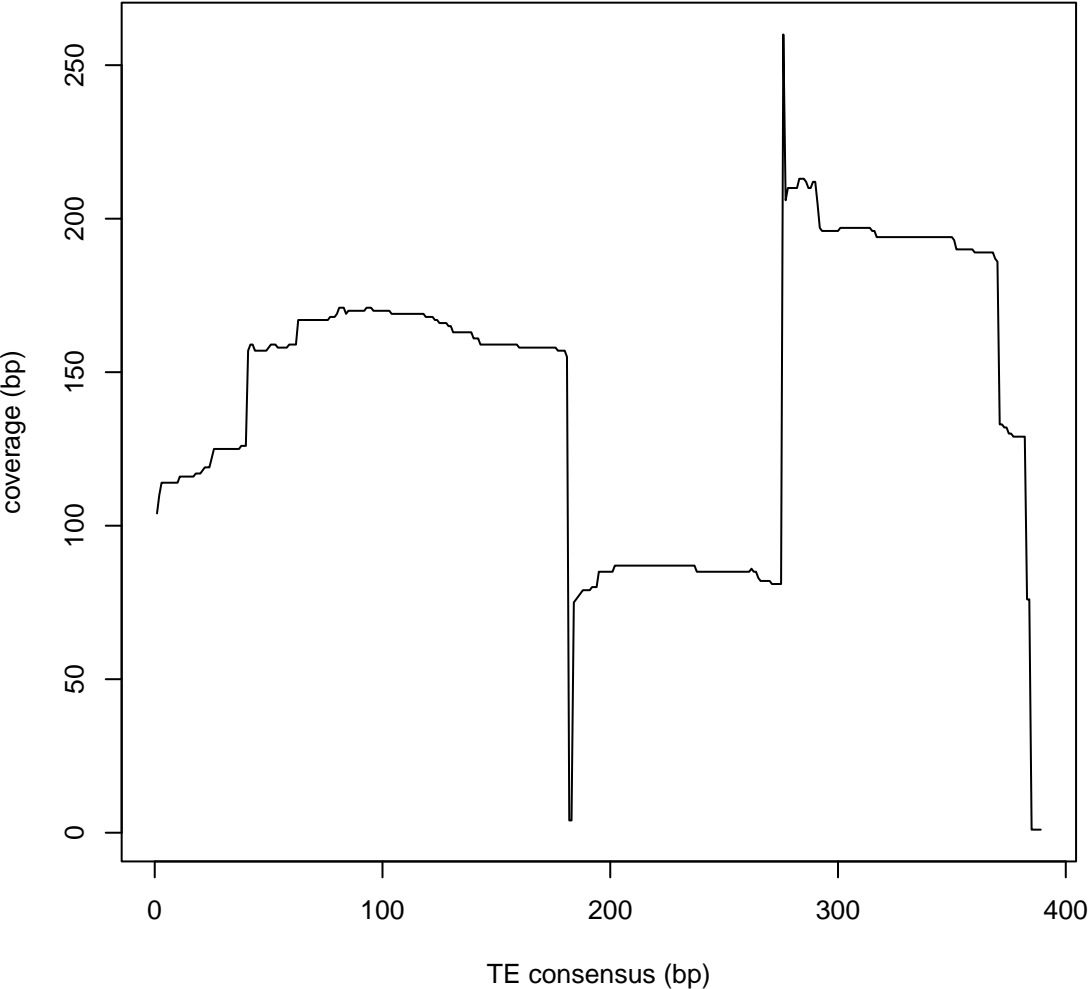


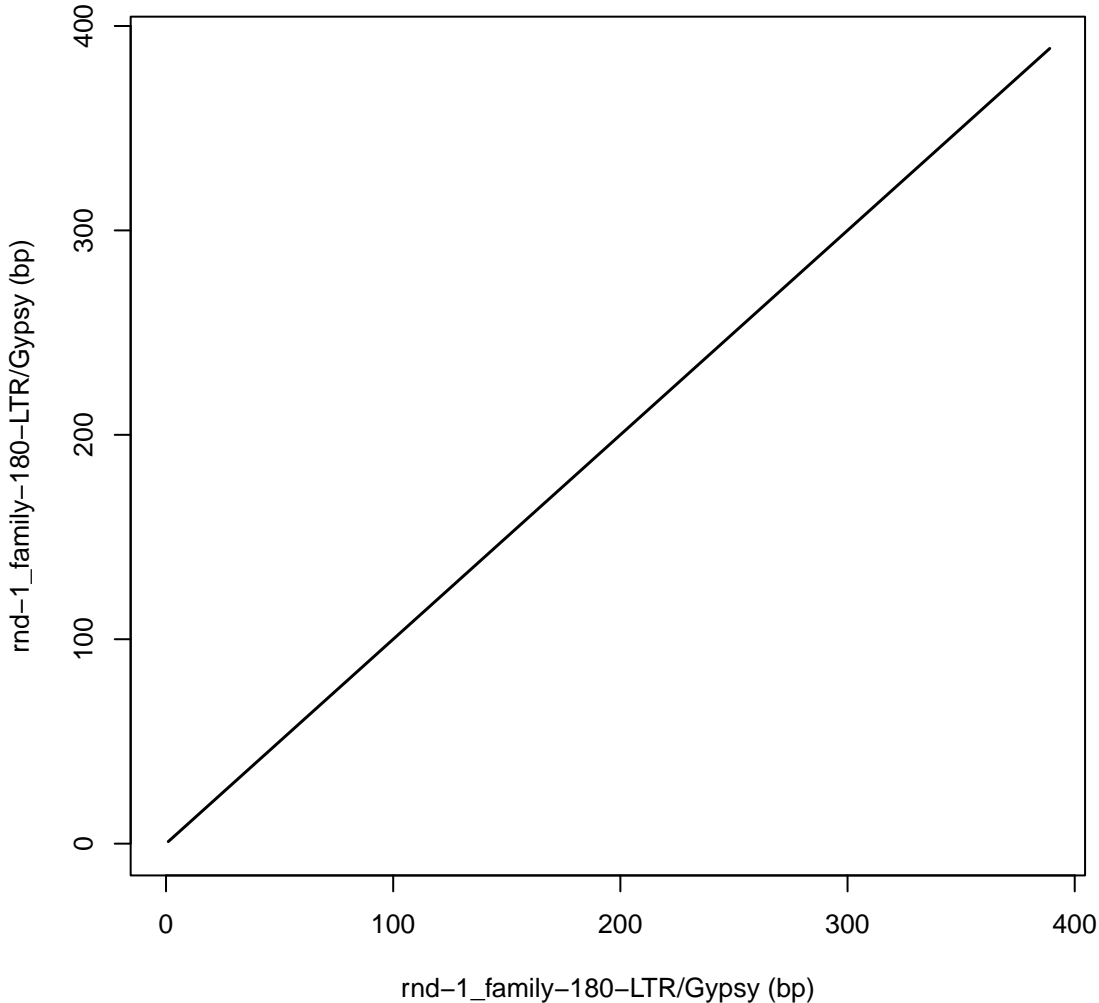
TE: rnd-1_family-180-LTR/Gypsy
consensus size: 389bp; fragments: 453; full length: 1 (>=350.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

