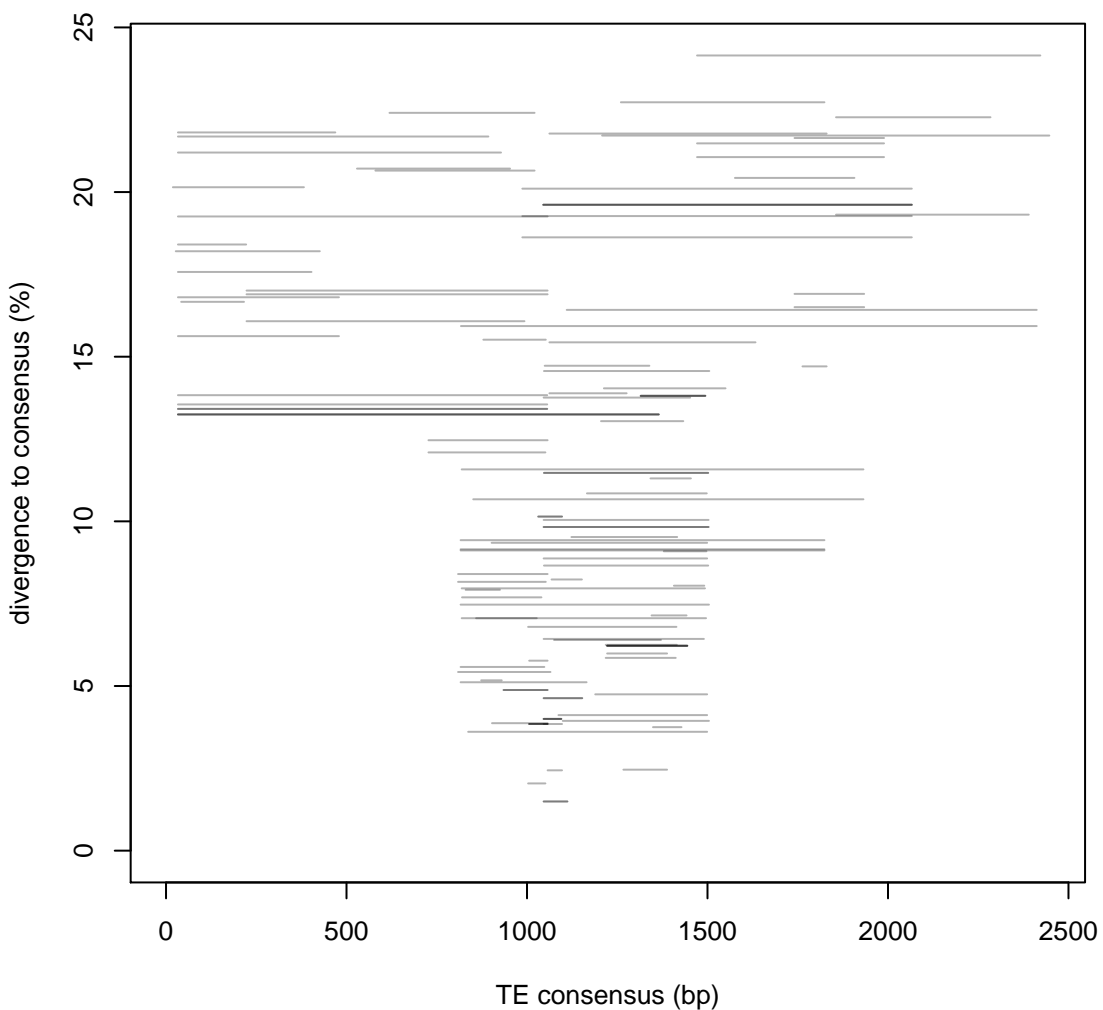
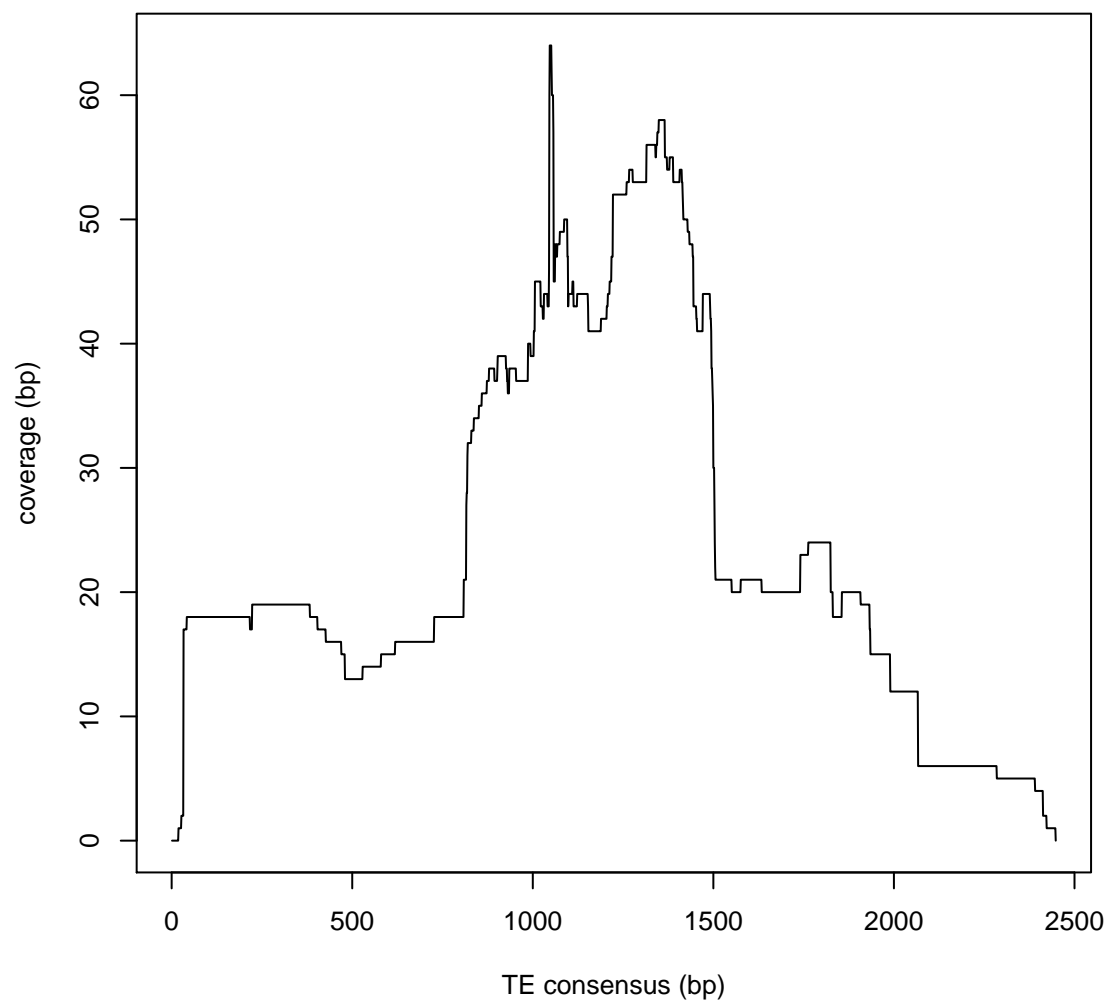


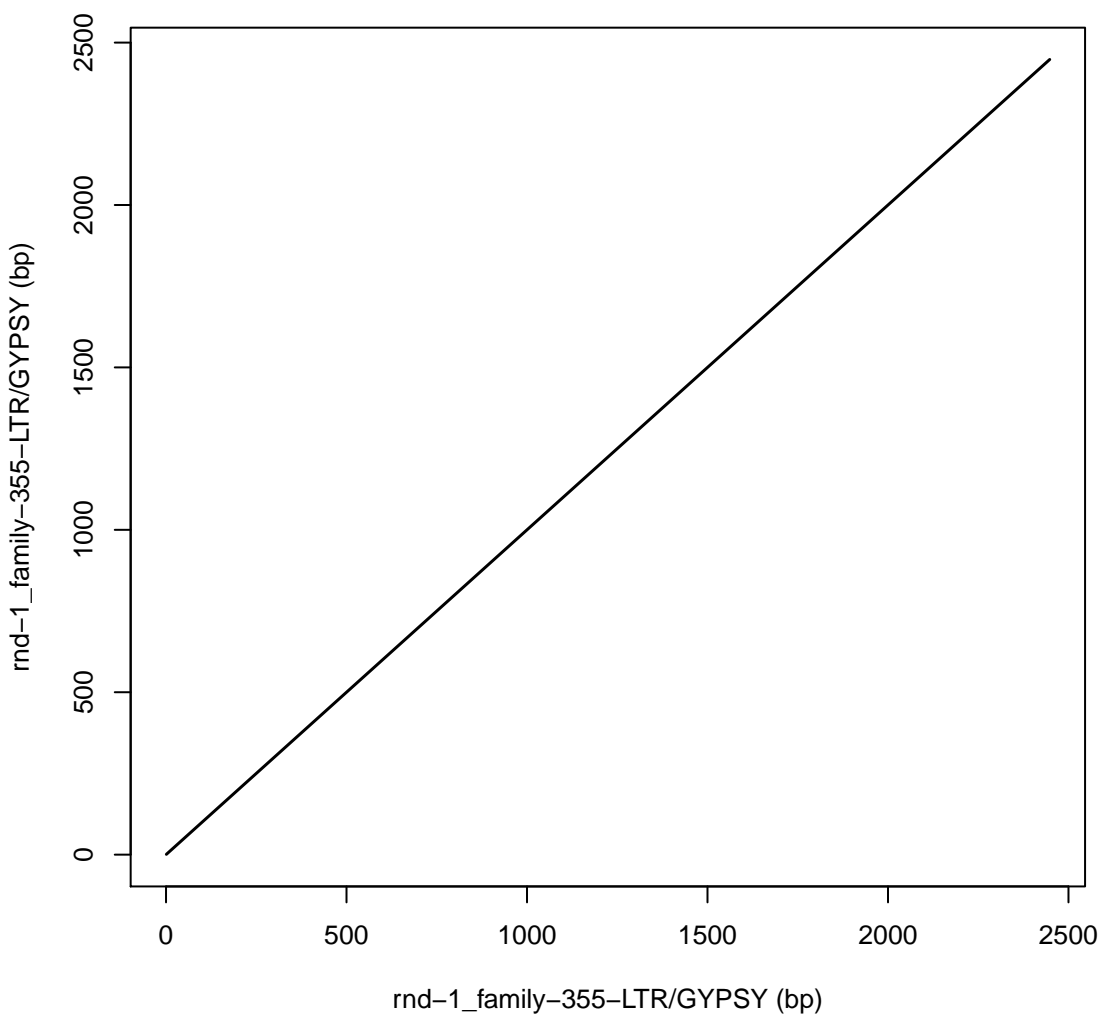
TE: rnd-1_family-355-LTR/GYPSY
consensus size: 2448bp; fragments: 125; full length: 0 (≥ 2203.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

