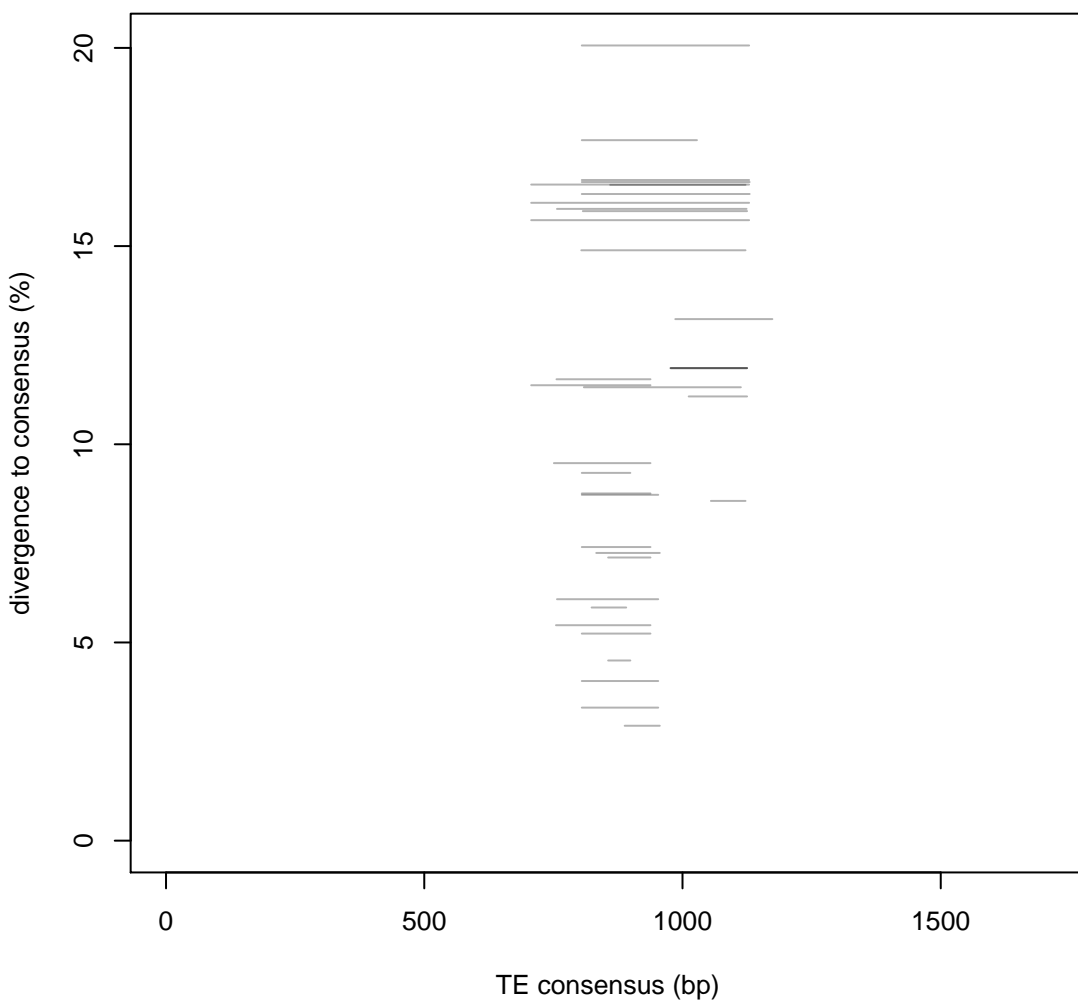
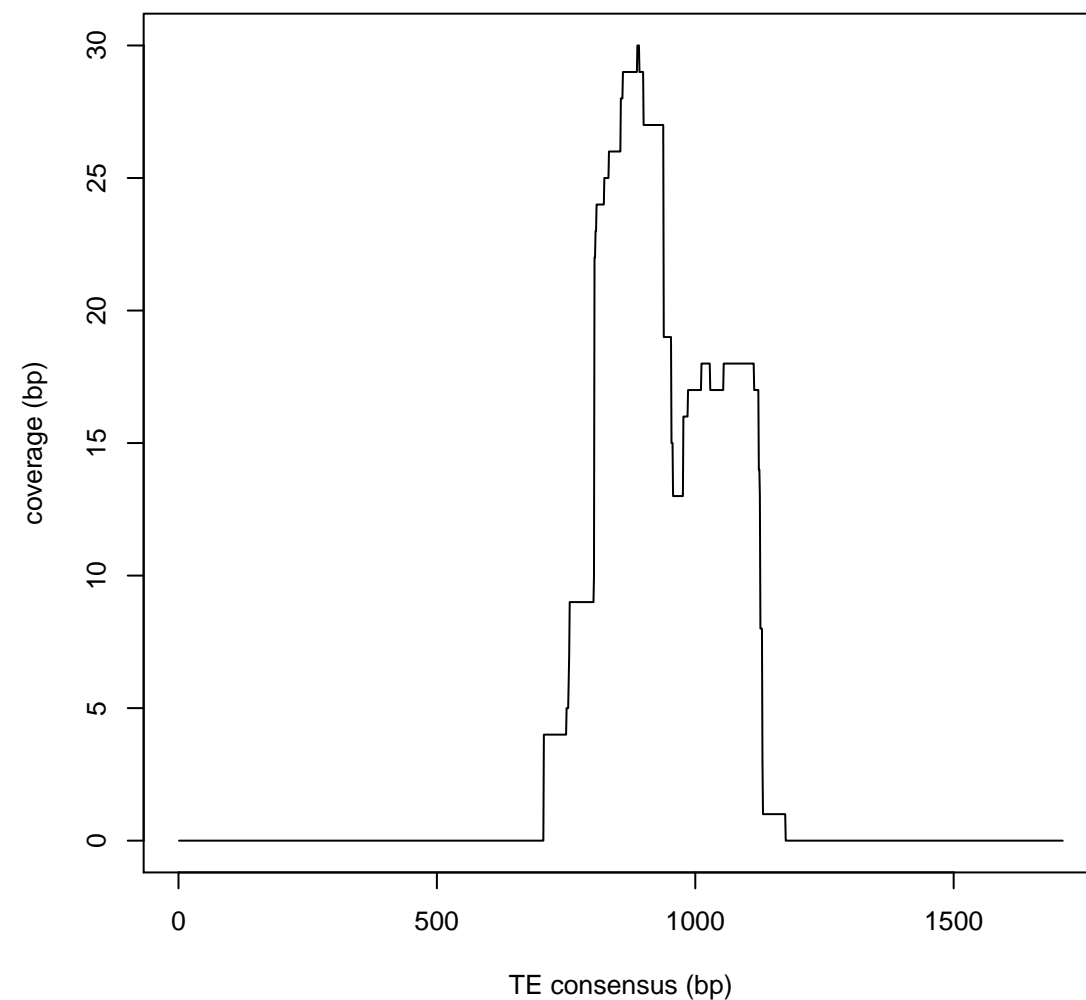


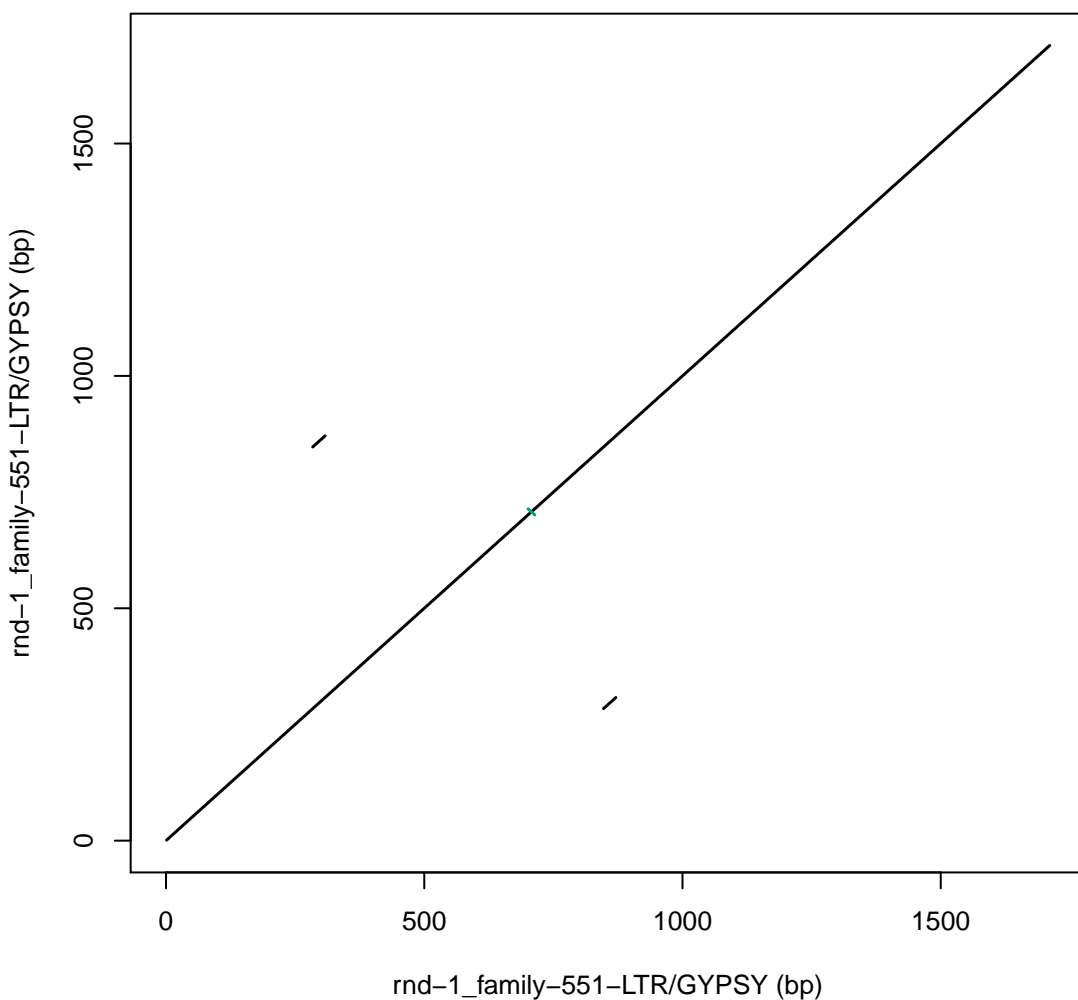
TE: rnd-1_family-551-LTR/GYPSY
consensus size: 1711bp; fragments: 36; full length: 0 (≥ 1539.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

