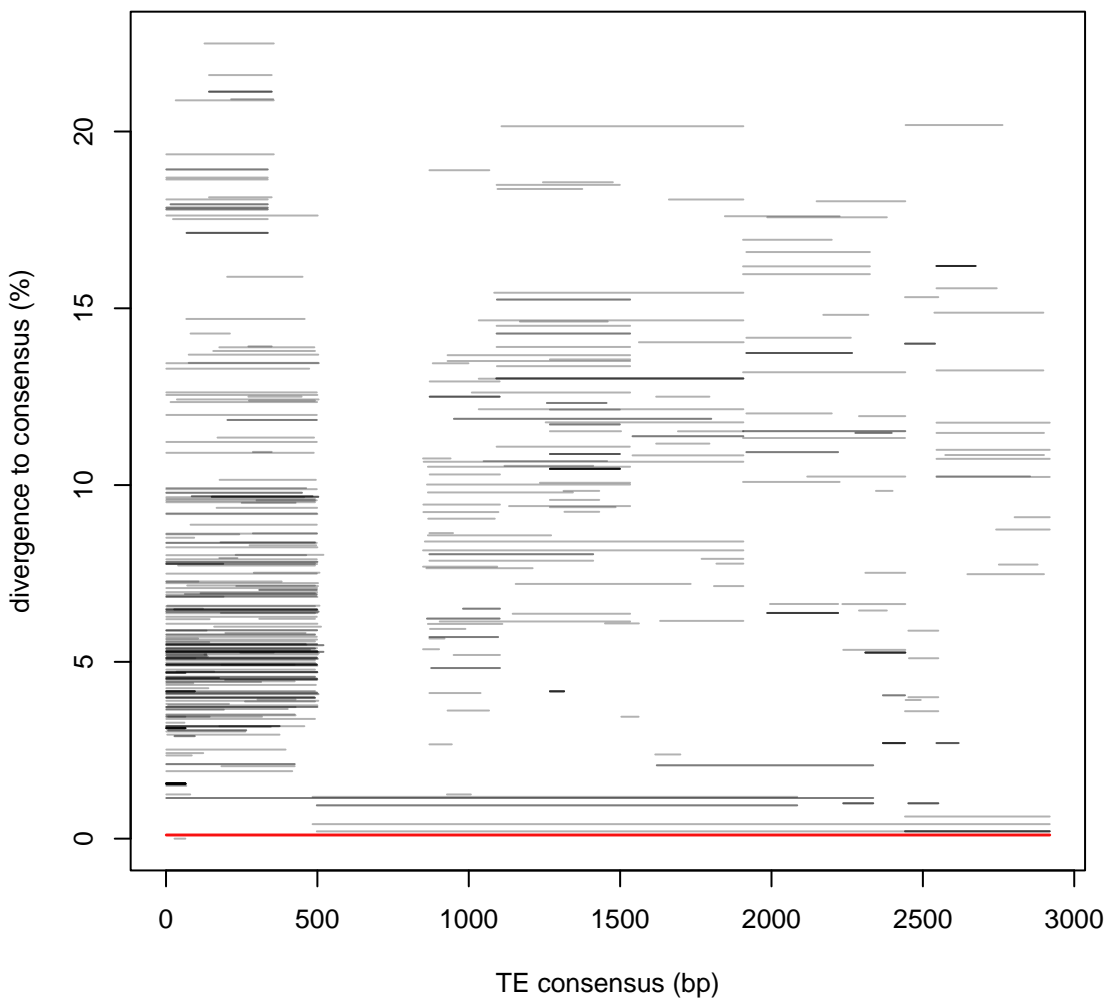
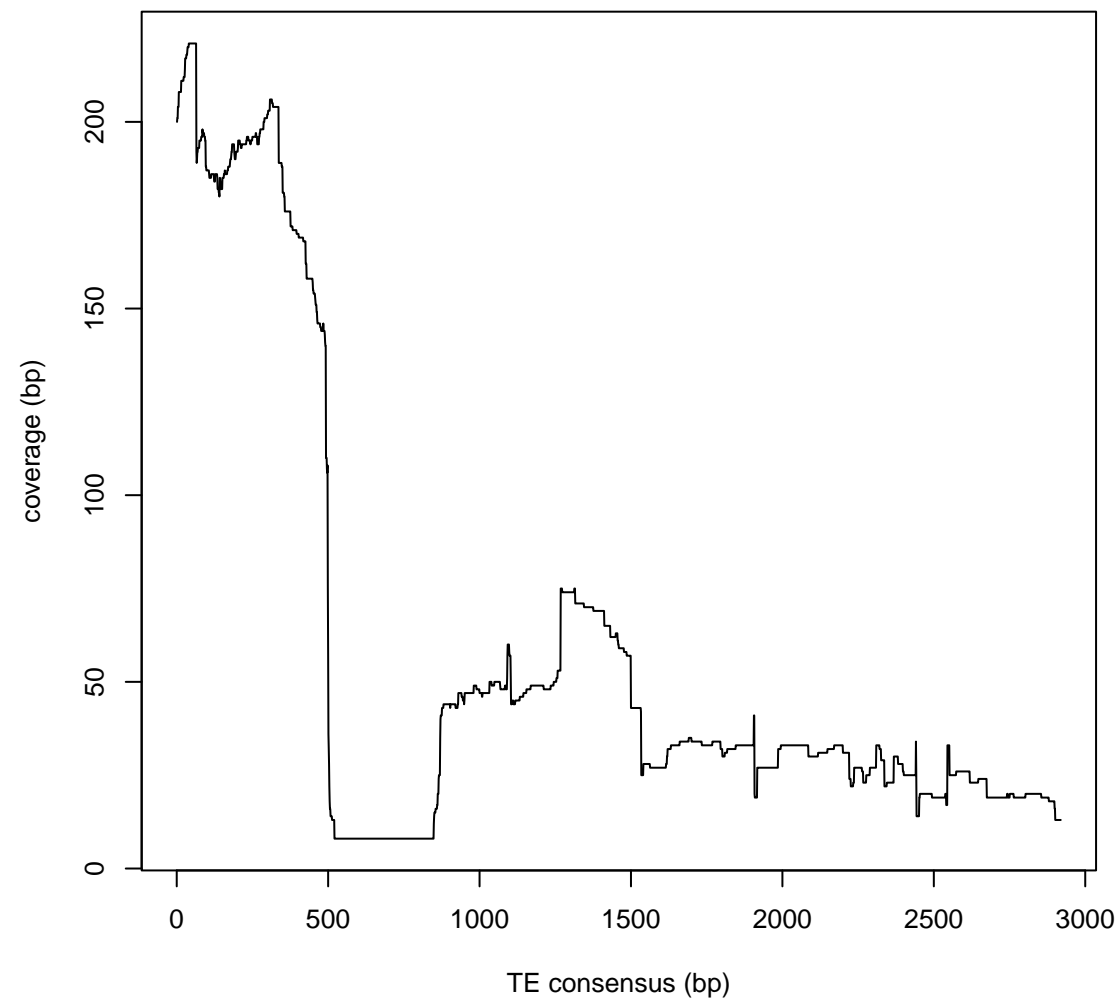


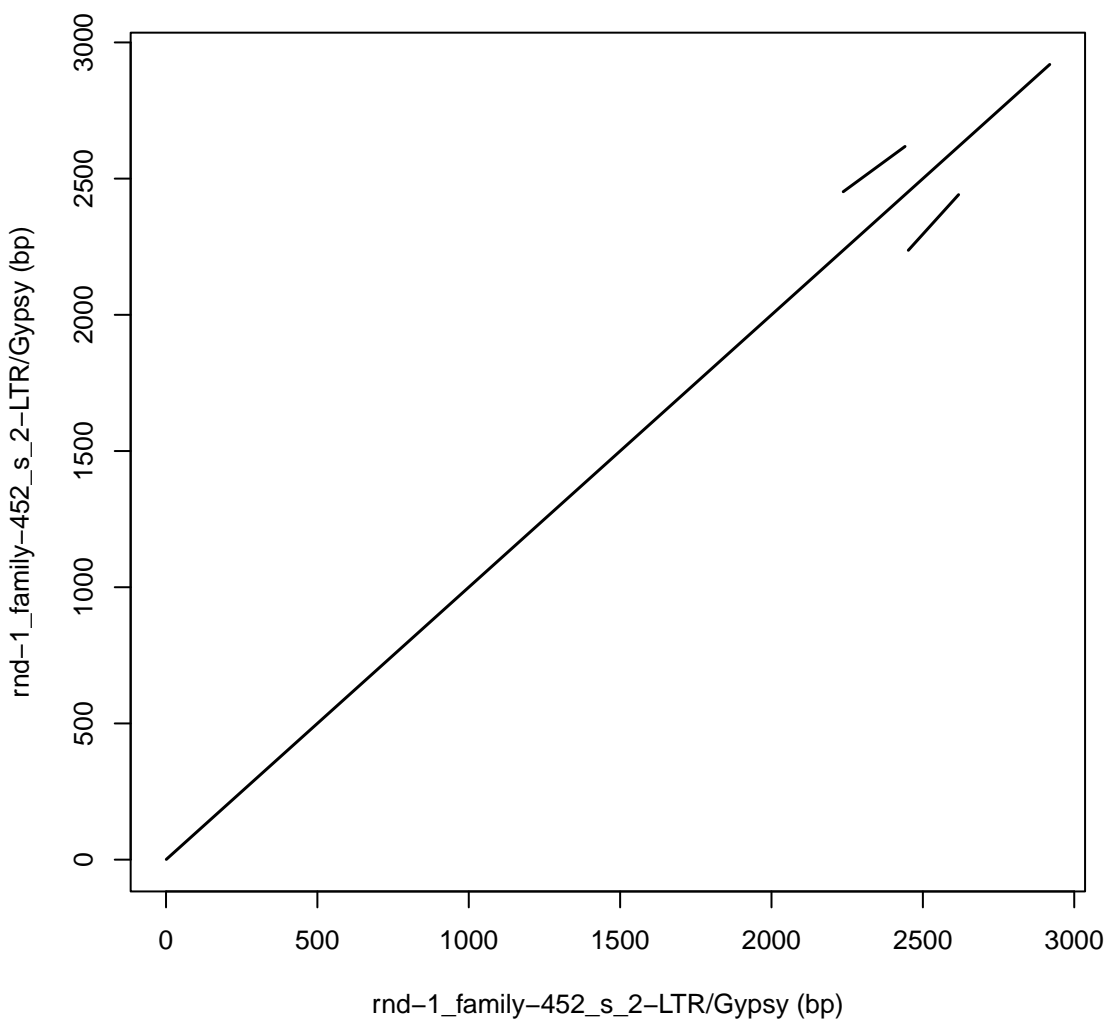
TE: rnd-1_family-452_s_2-LTR/Gypsy
consensus size: 2919bp; fragments: 489; full length: 1 (>=2627.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

