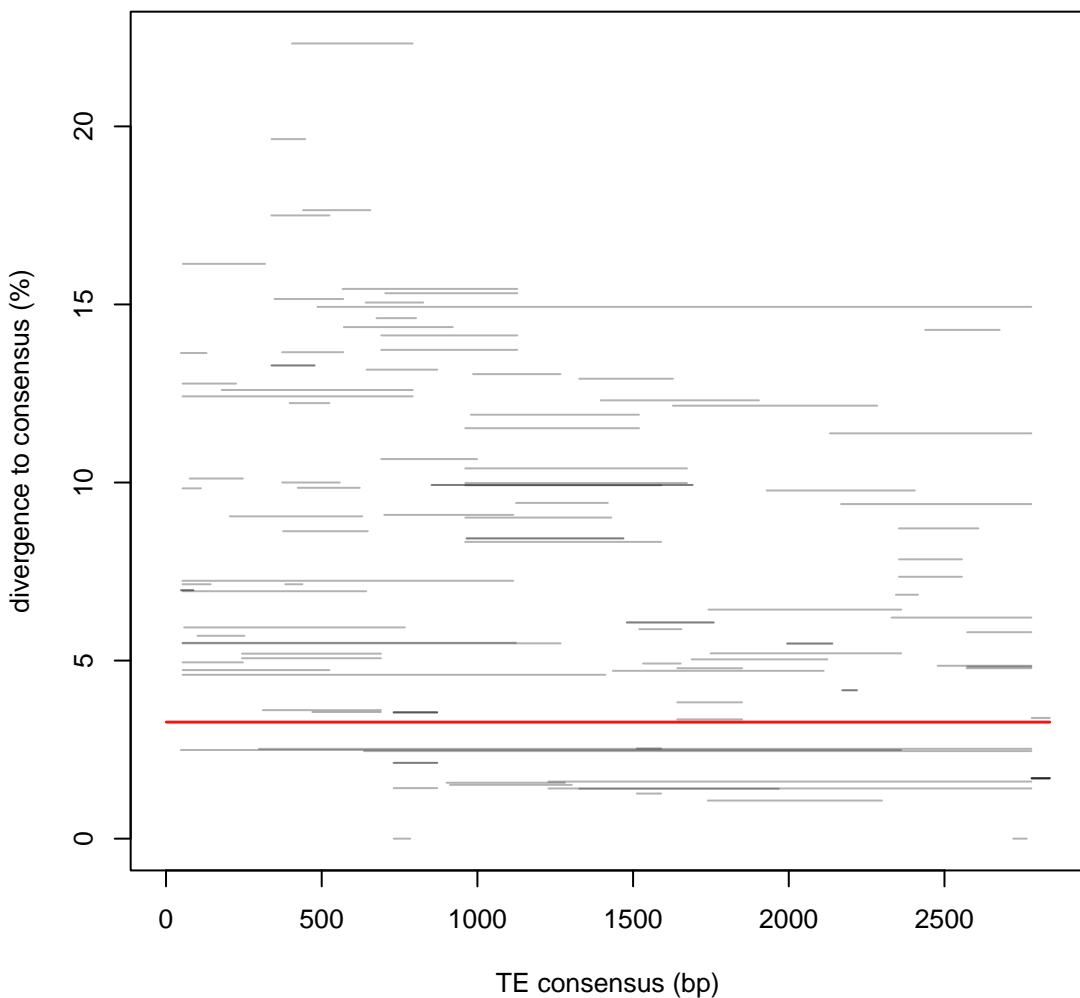
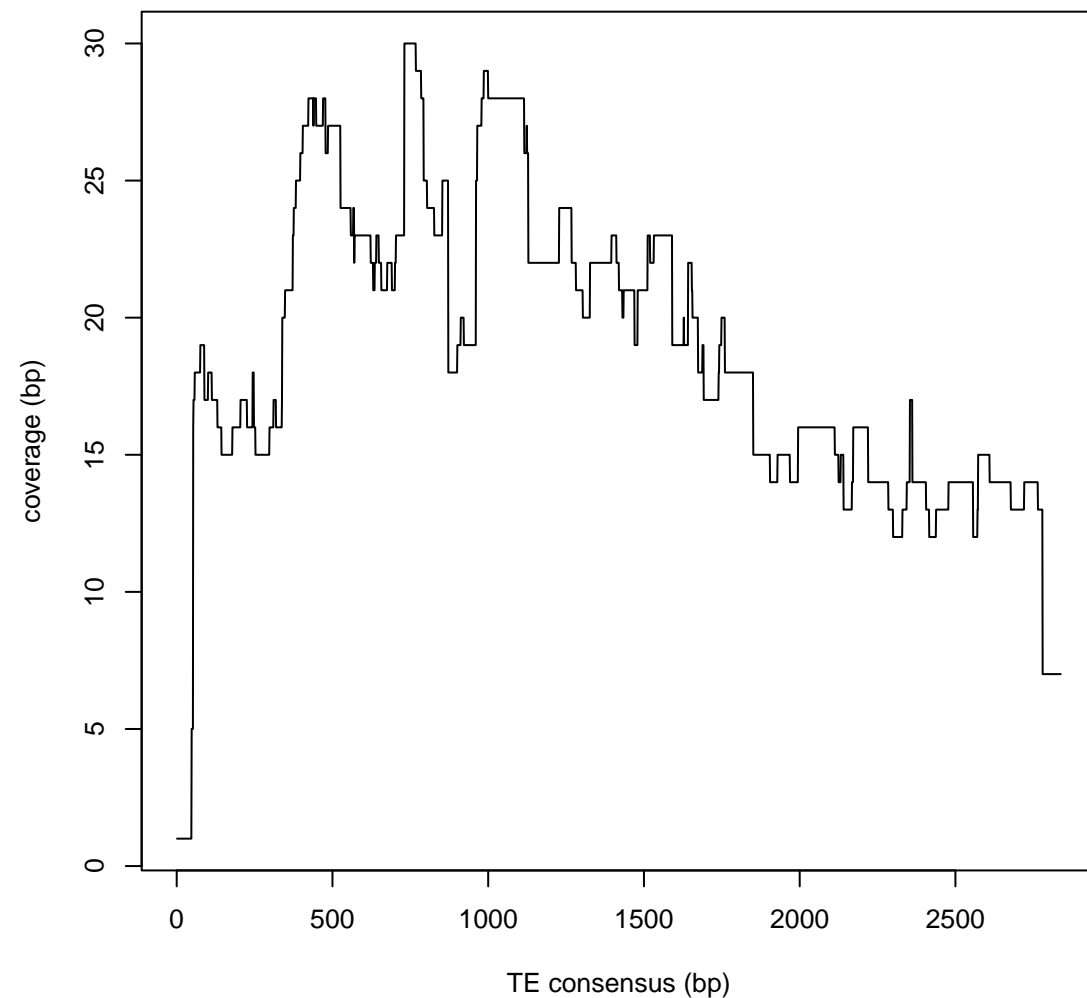


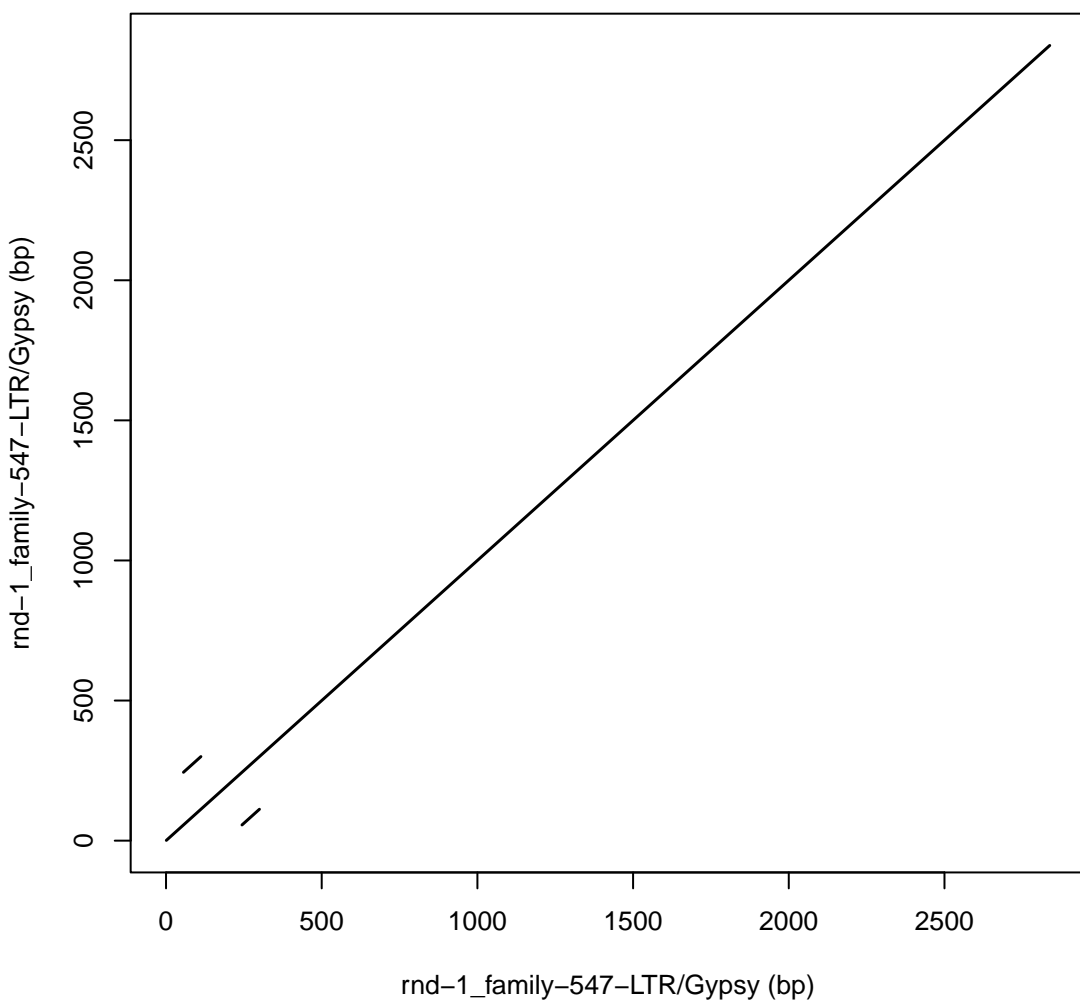
TE: rnd-1_family-547-LTR/Gypsy
consensus size: 2838bp; fragments: 118; full length: 1 (>=2554.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

