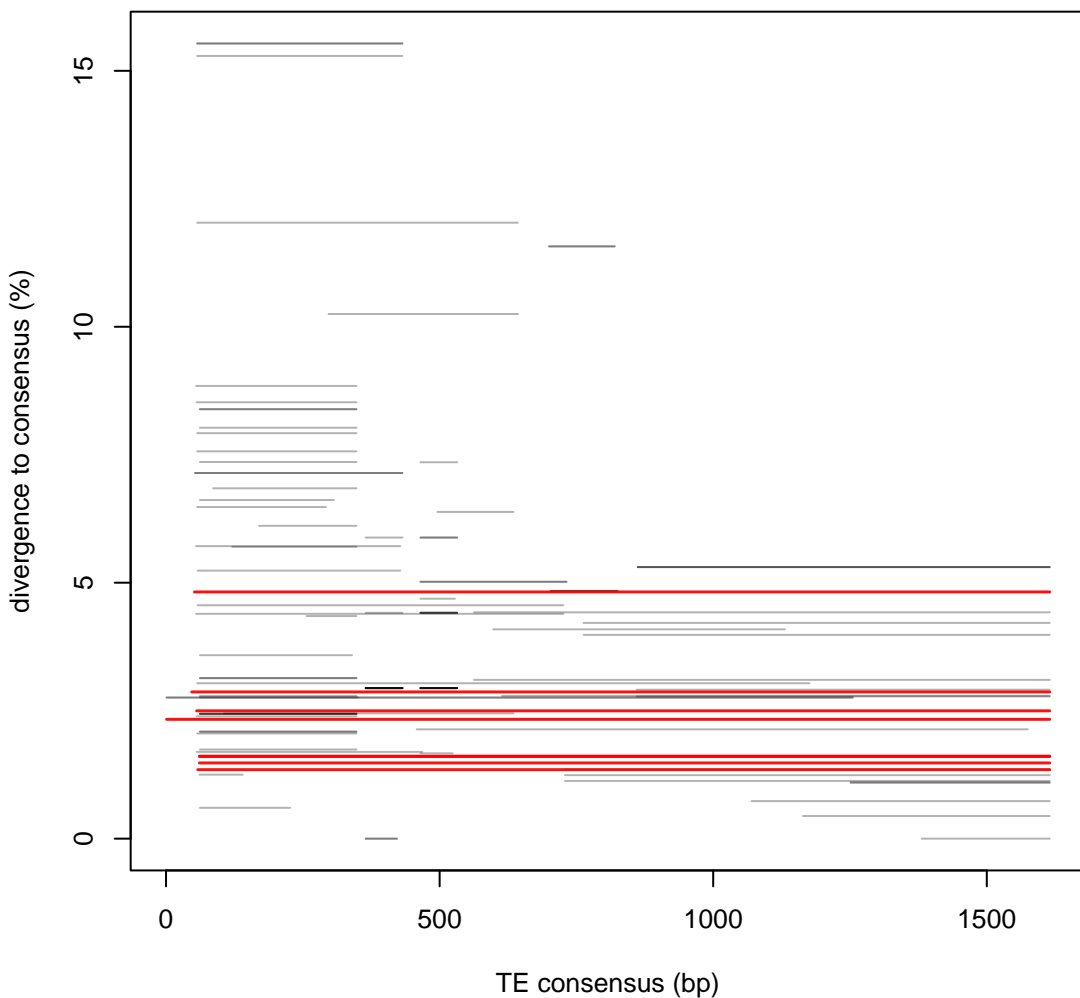
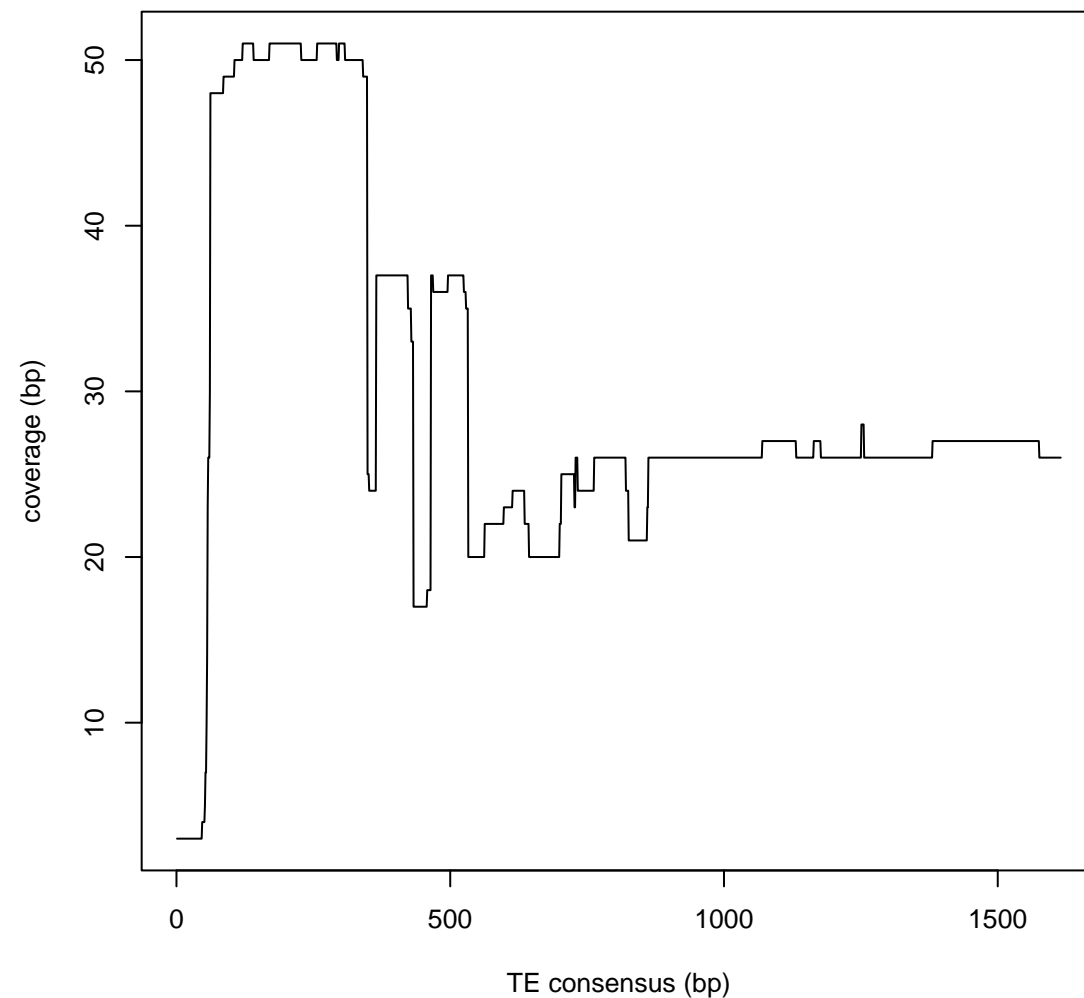


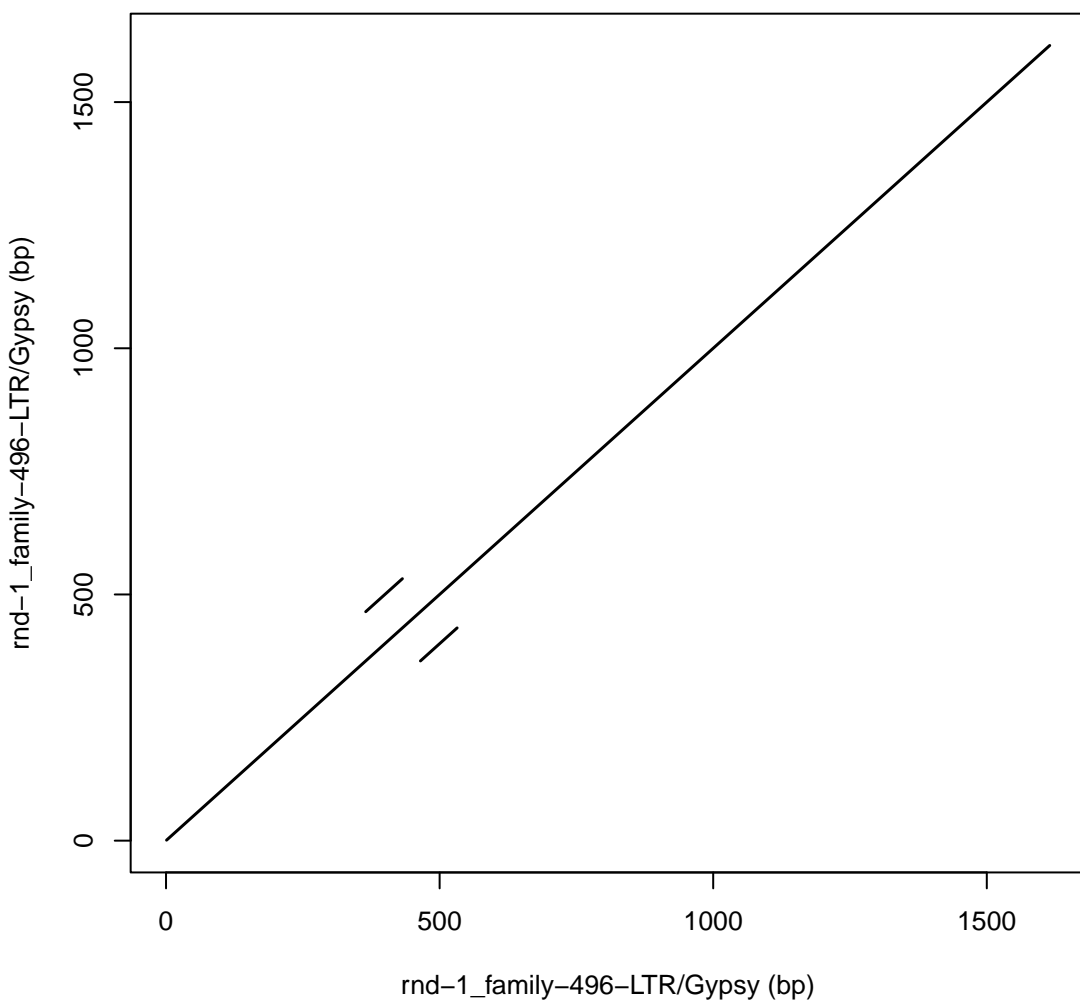
TE: rnd-1_family-496-LTR/Gypsy
consensus size: 1615bp; fragments: 112; full length: 8 (>=1453.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

