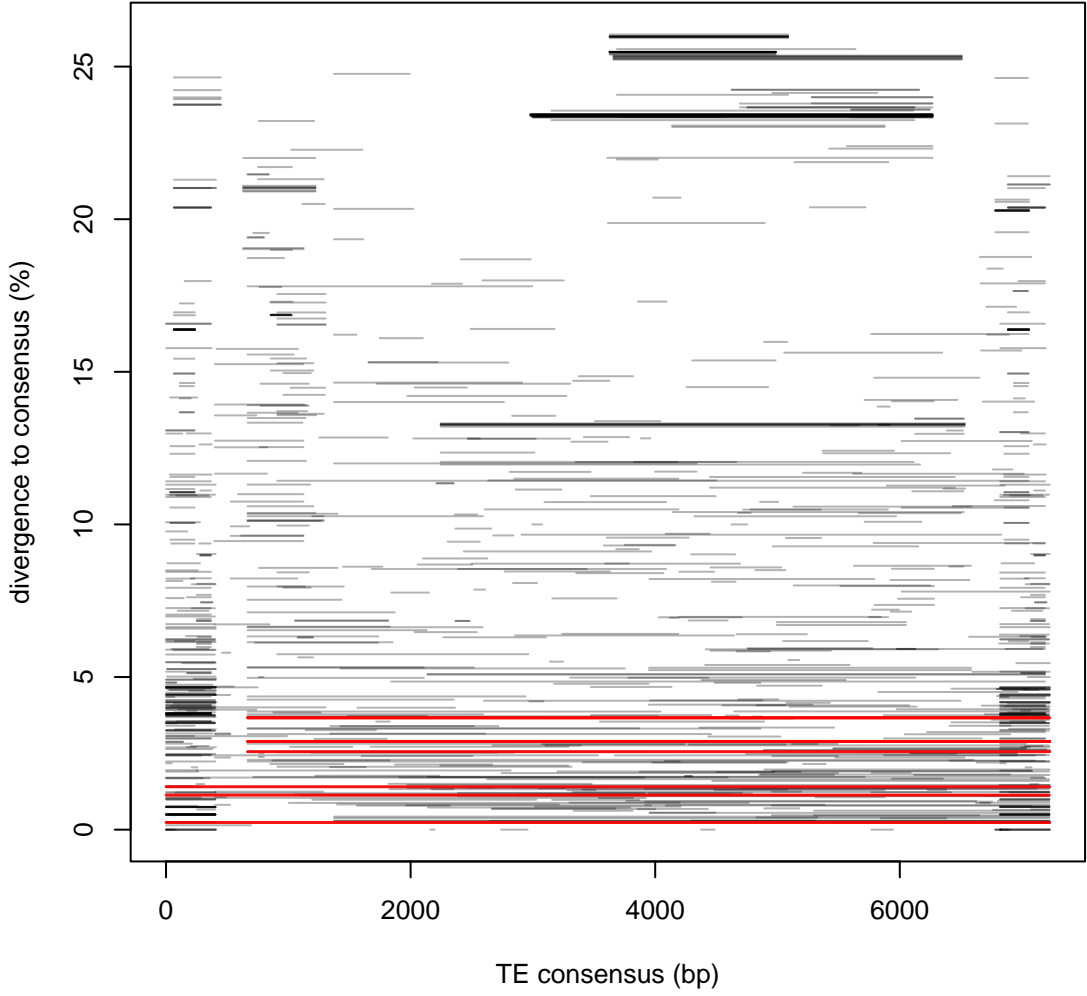
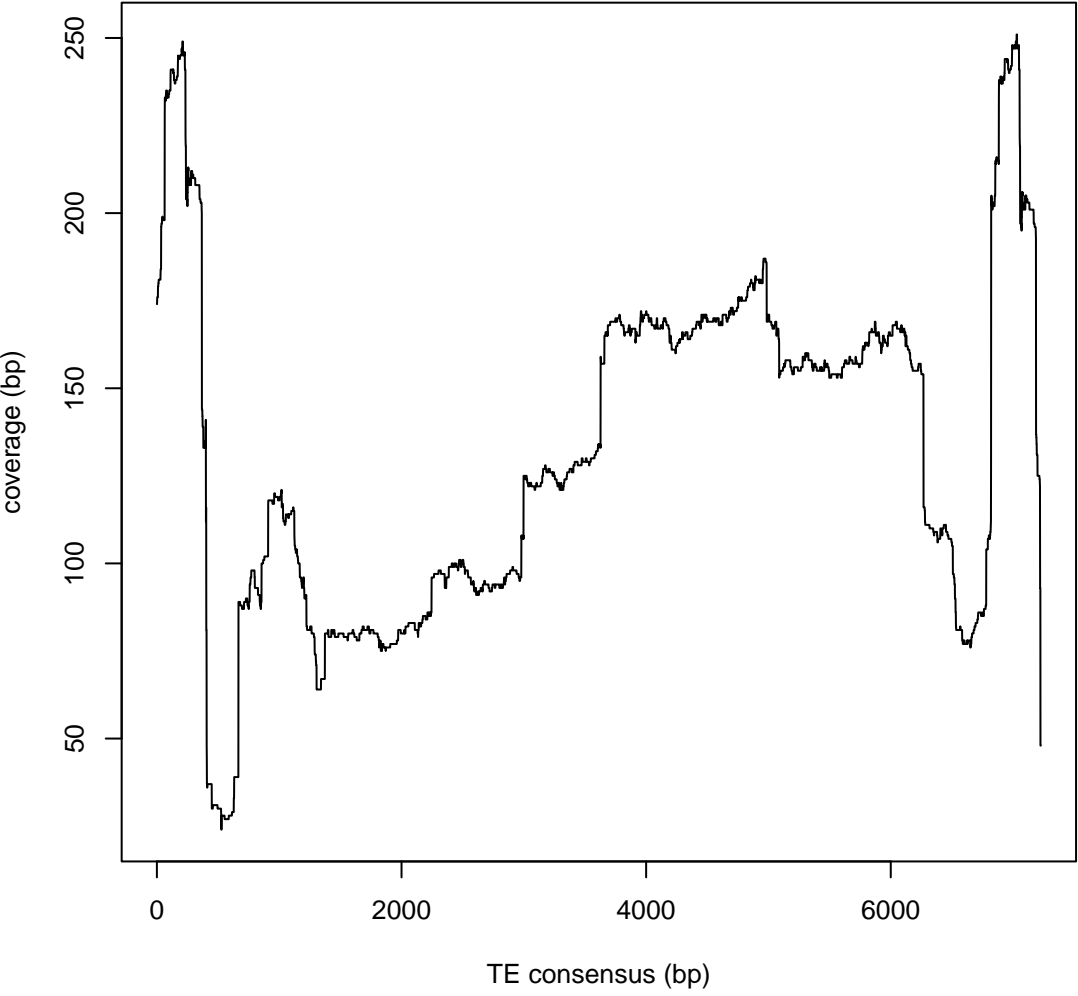


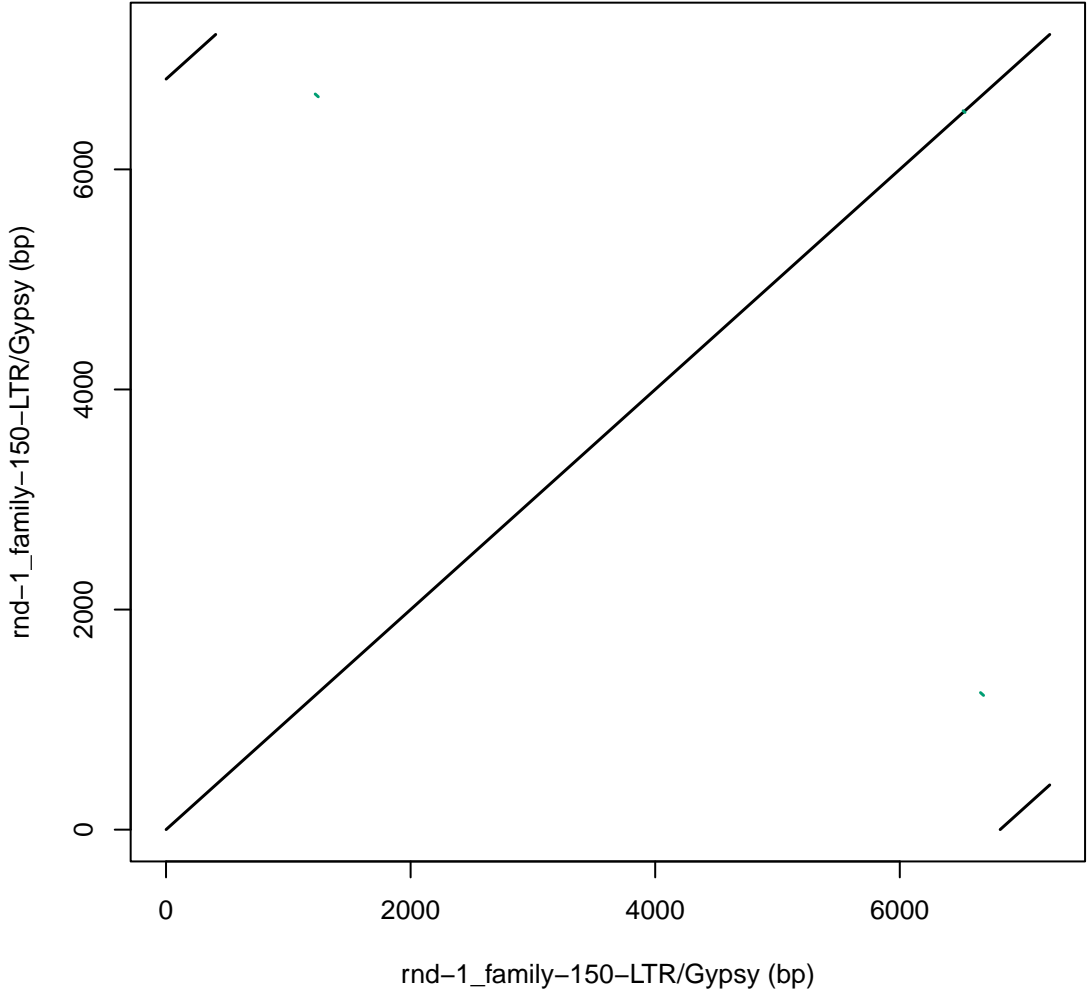
TE: rnd-1_family-150-LTR/Gypsy
consensus size: 7226bp; fragments: 1065; full length: 8 (>=6503.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

