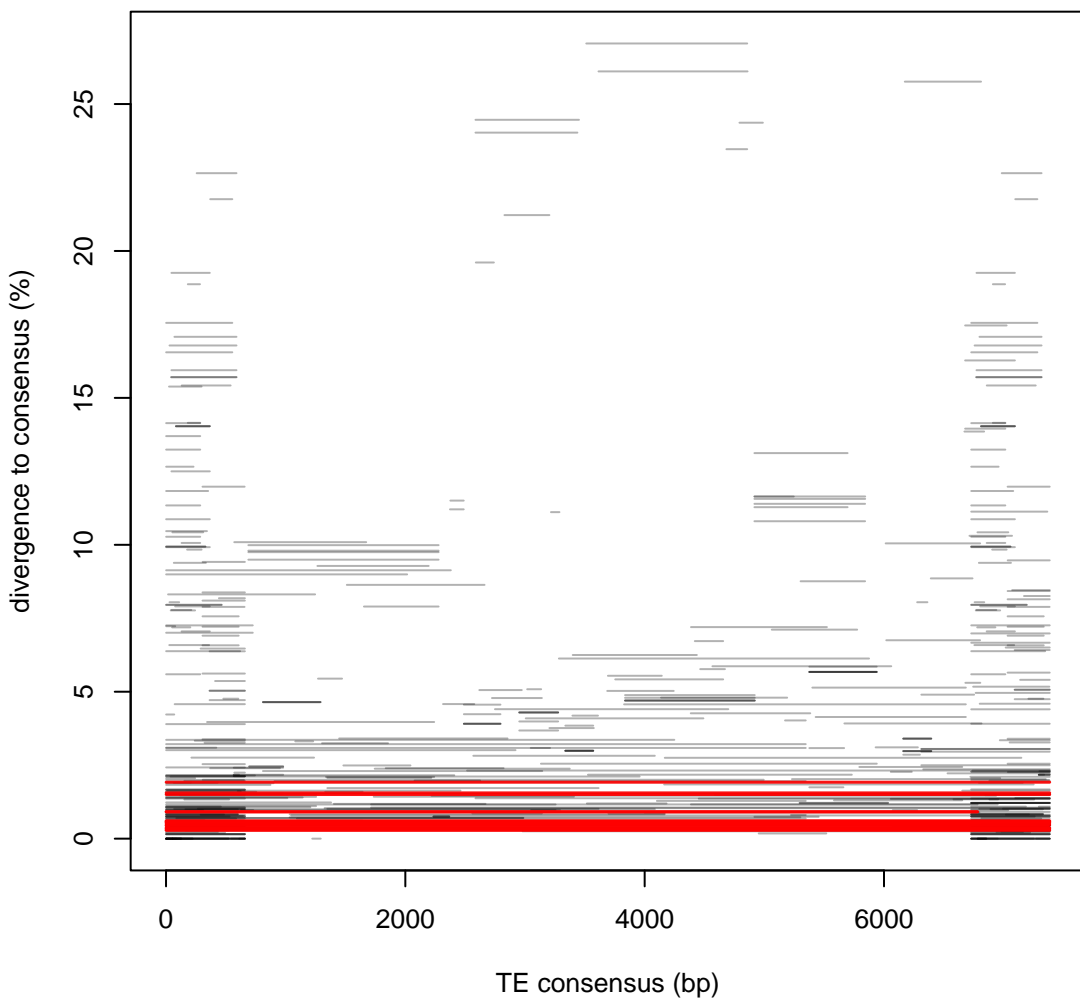
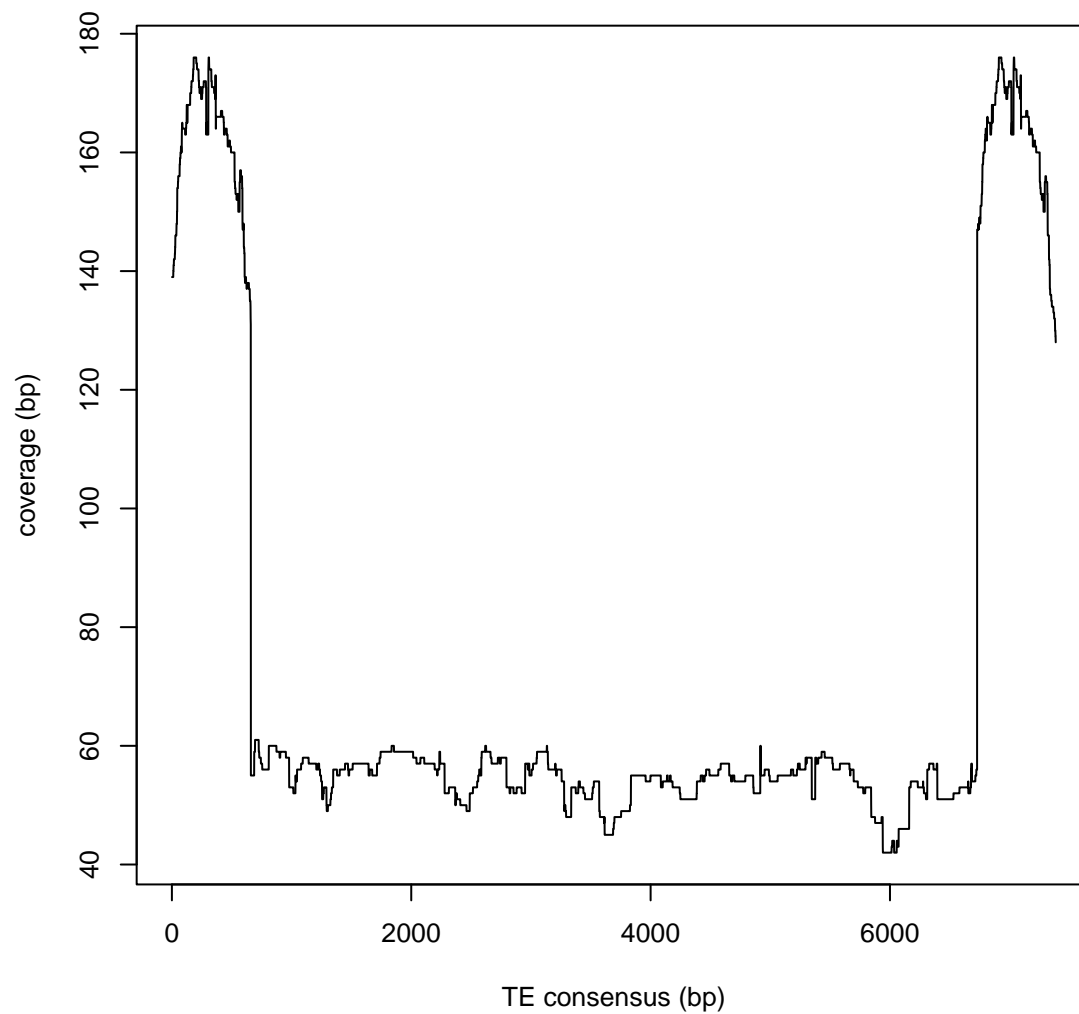


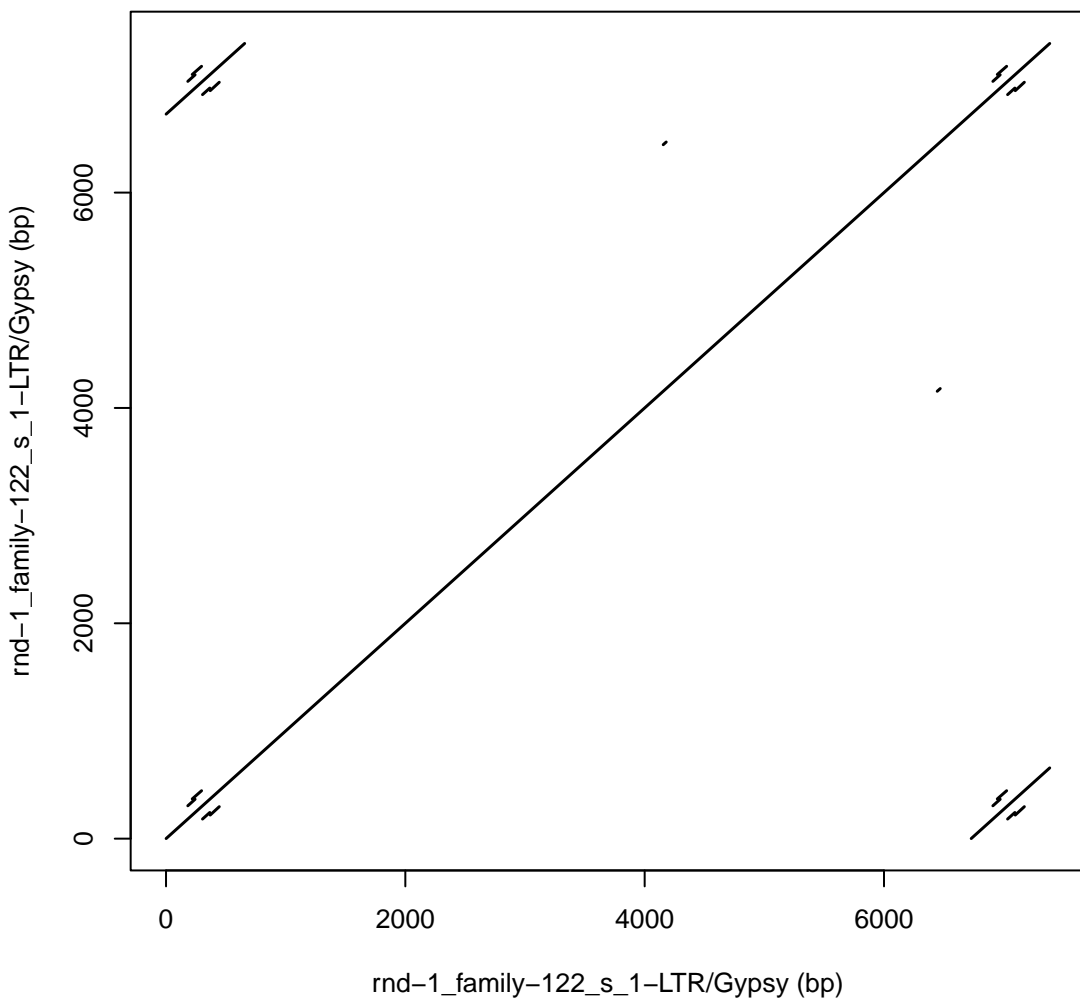
TE: rnd-1_family-122_s_1-LTR/Gypsy
consensus size: 7385bp; fragments: 578; full length: 16 (>=6646.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

