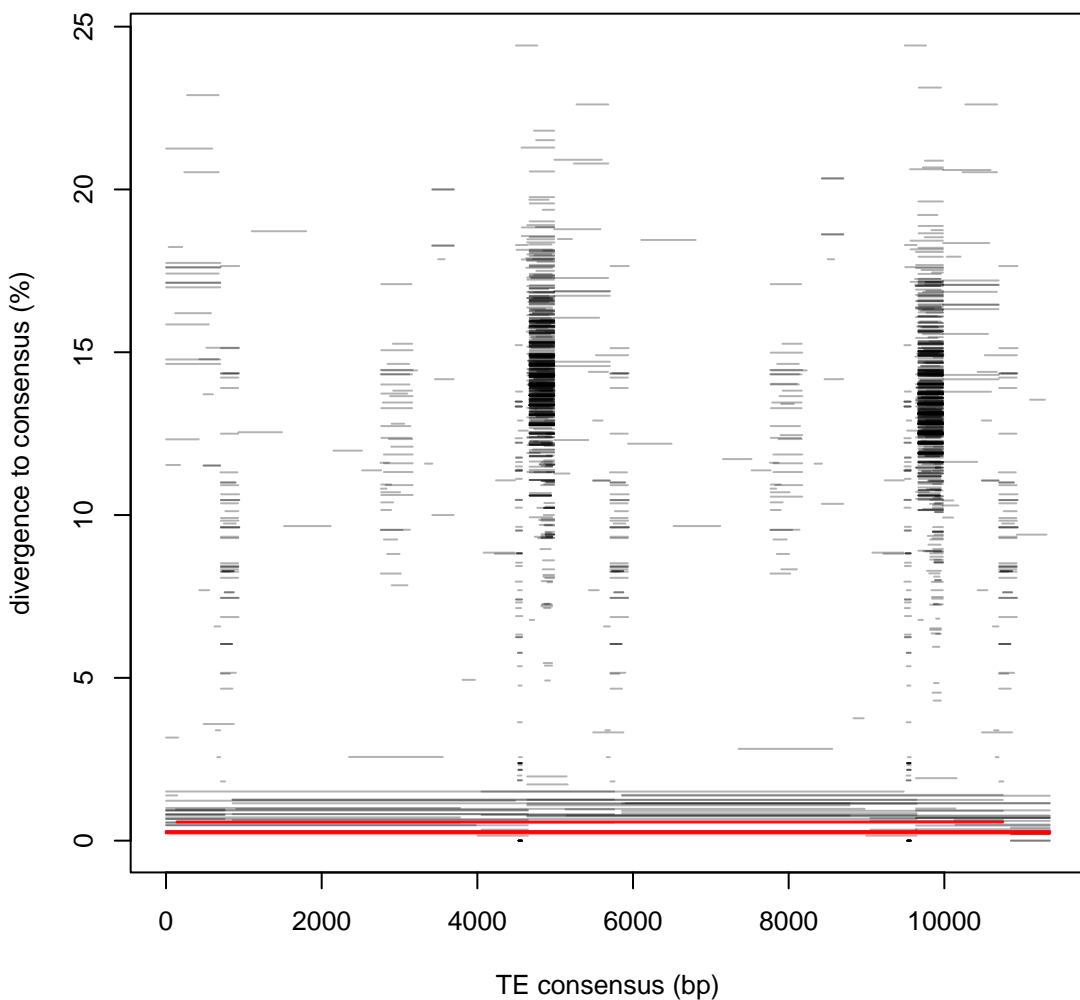
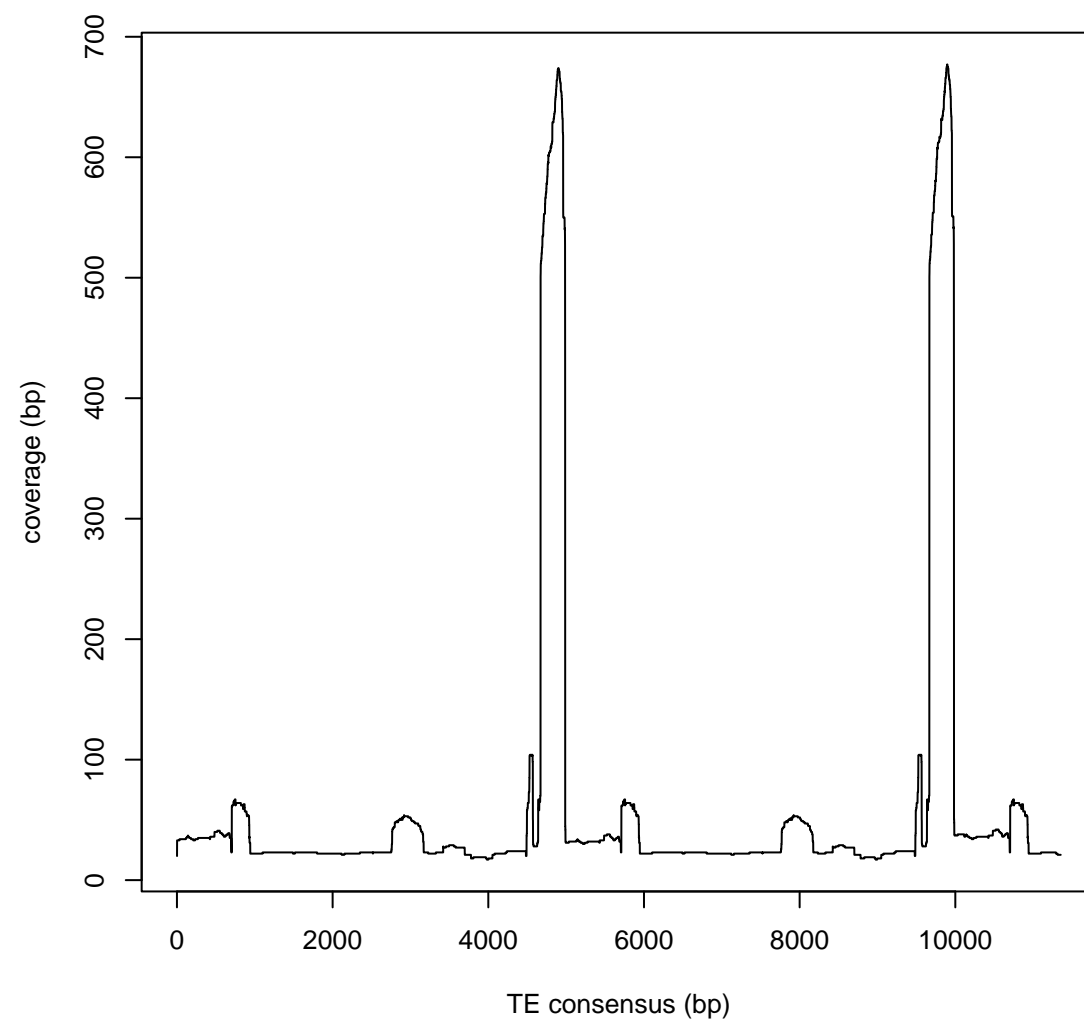


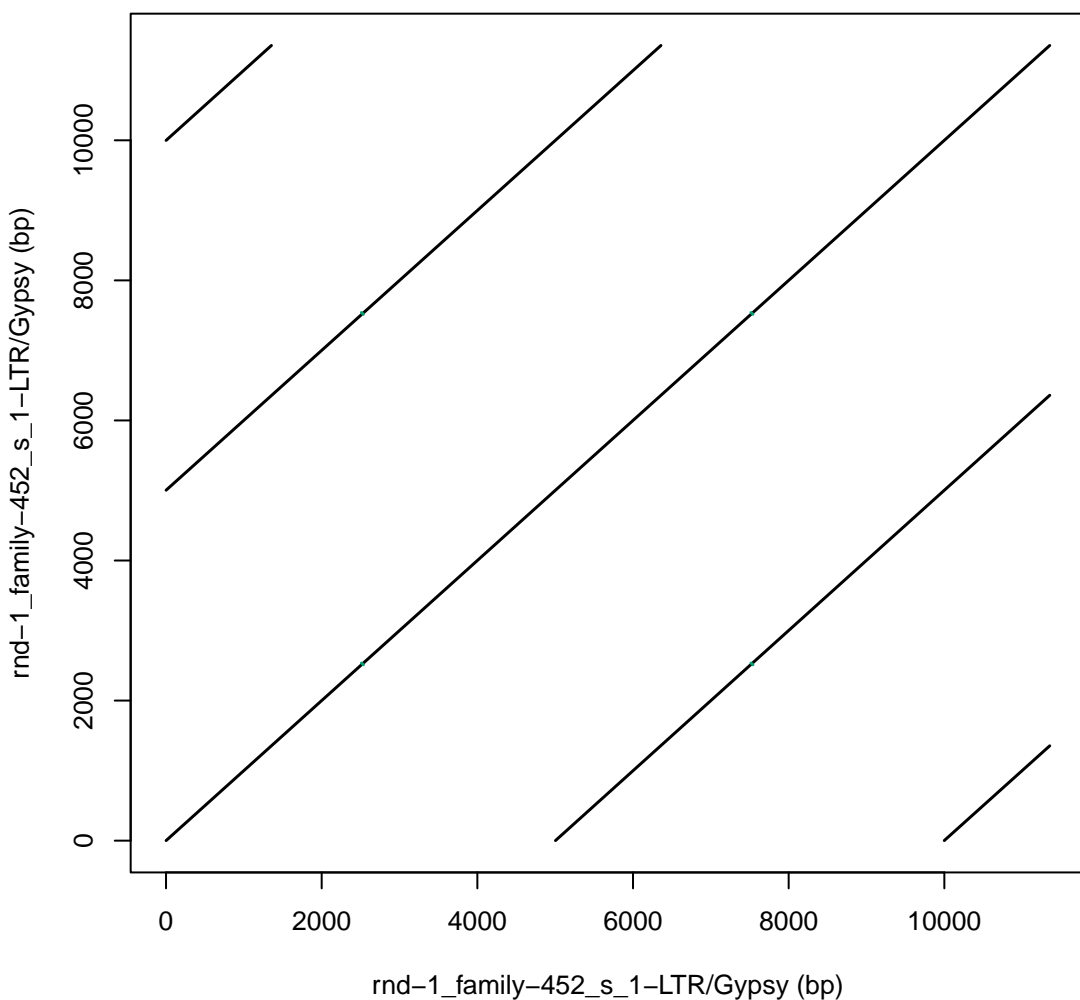
TE: rnd-1_family-452_s_1-LTR/Gypsy
consensus size: 11354bp; fragments: 1893; full length: 3 (>=10218.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

