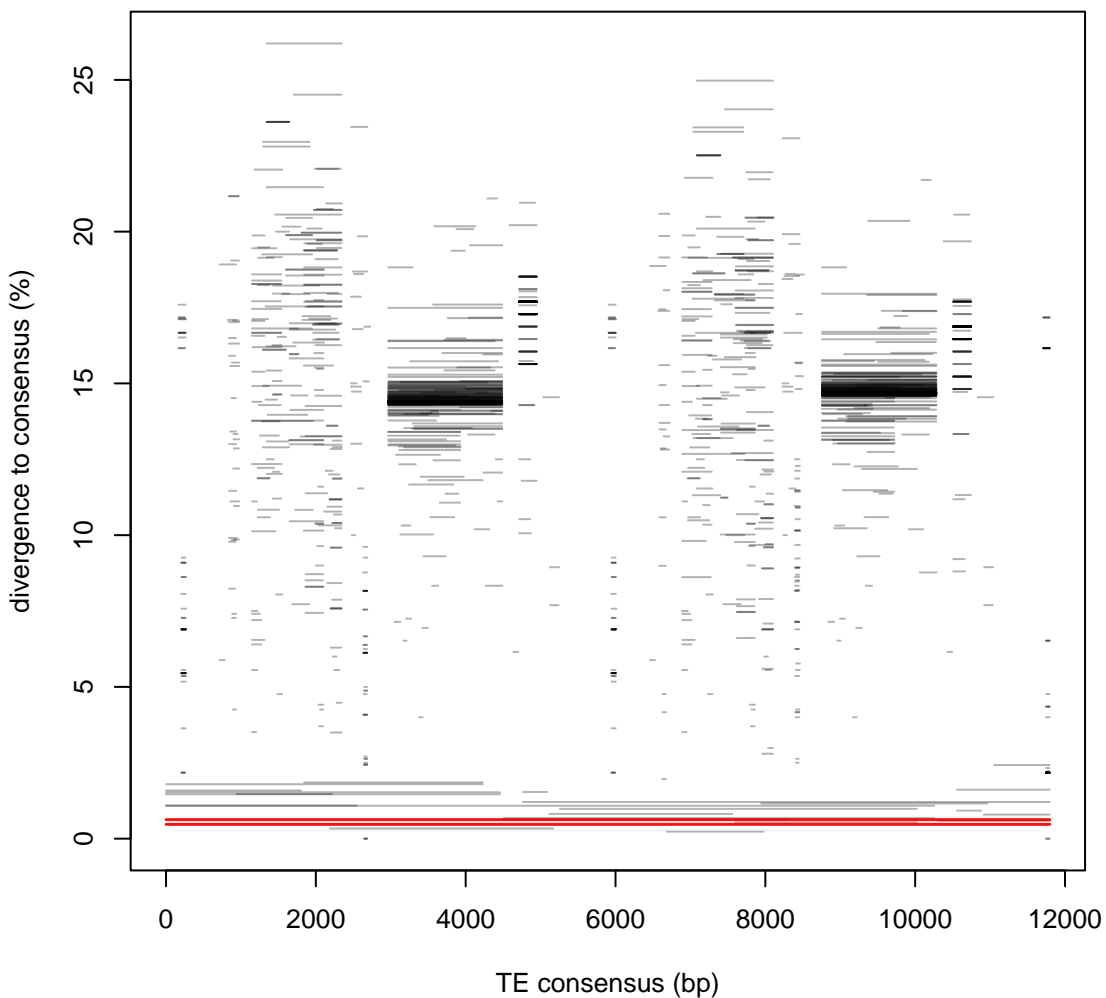
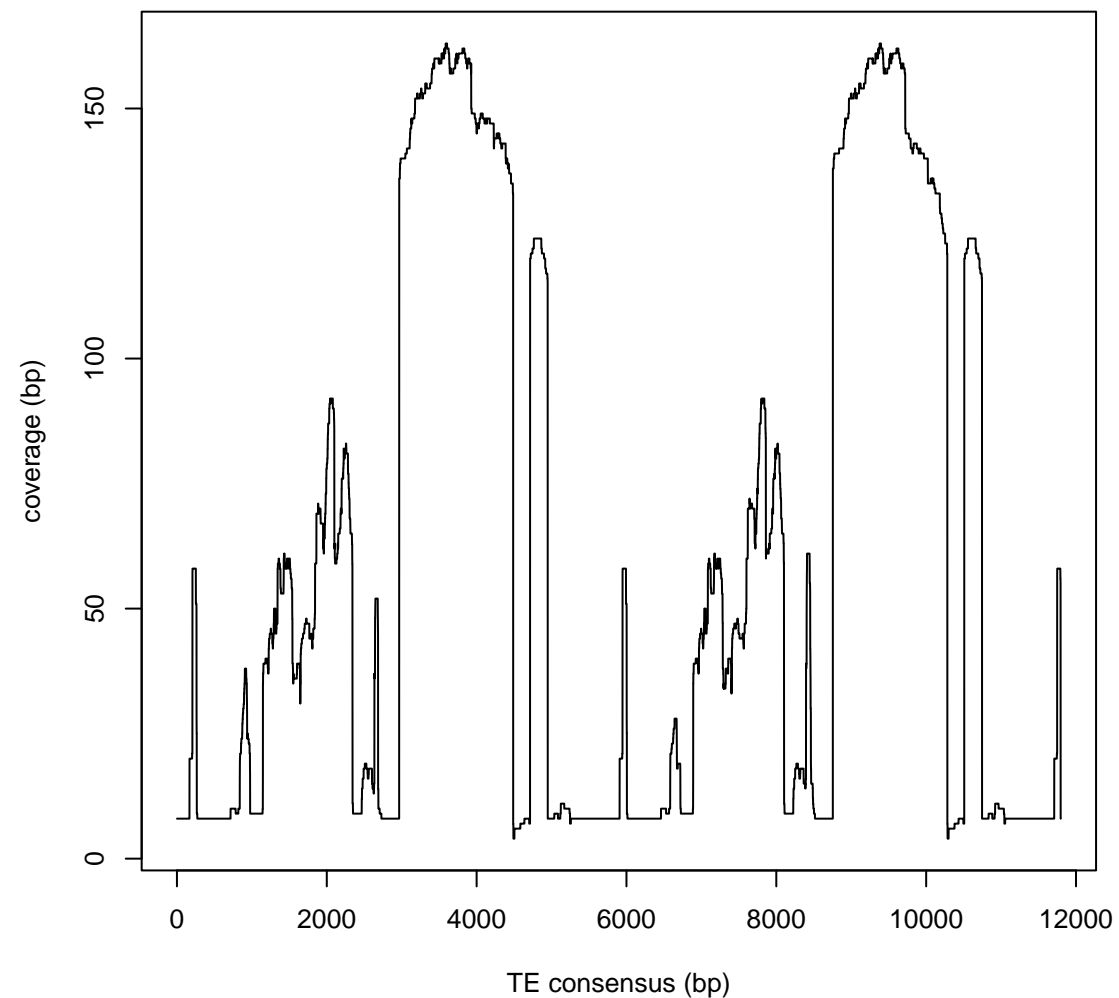


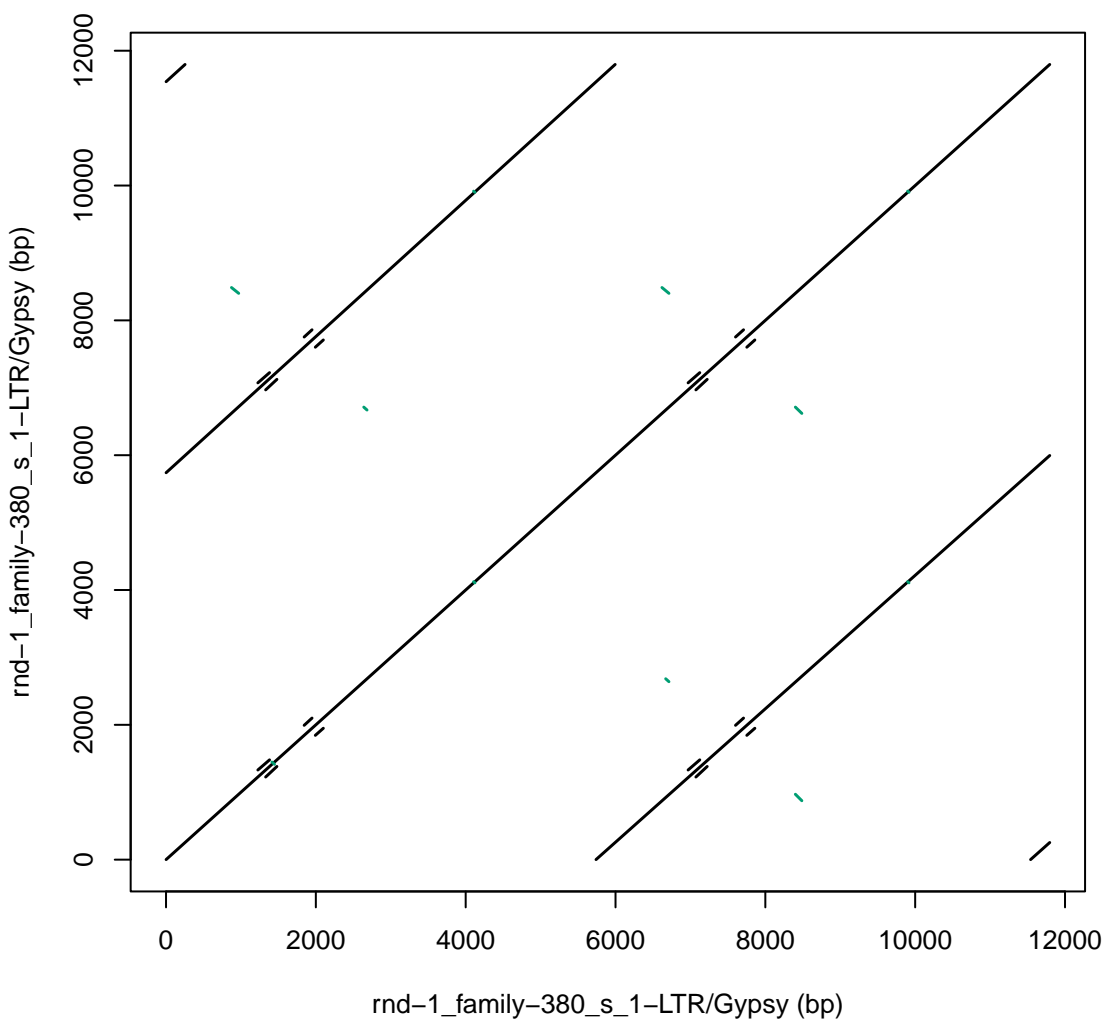
TE: rnd-1_family-380_s_1-LTR/Gypsy
consensus size: 11796bp; fragments: 1355; full length: 2 (>=10616.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

