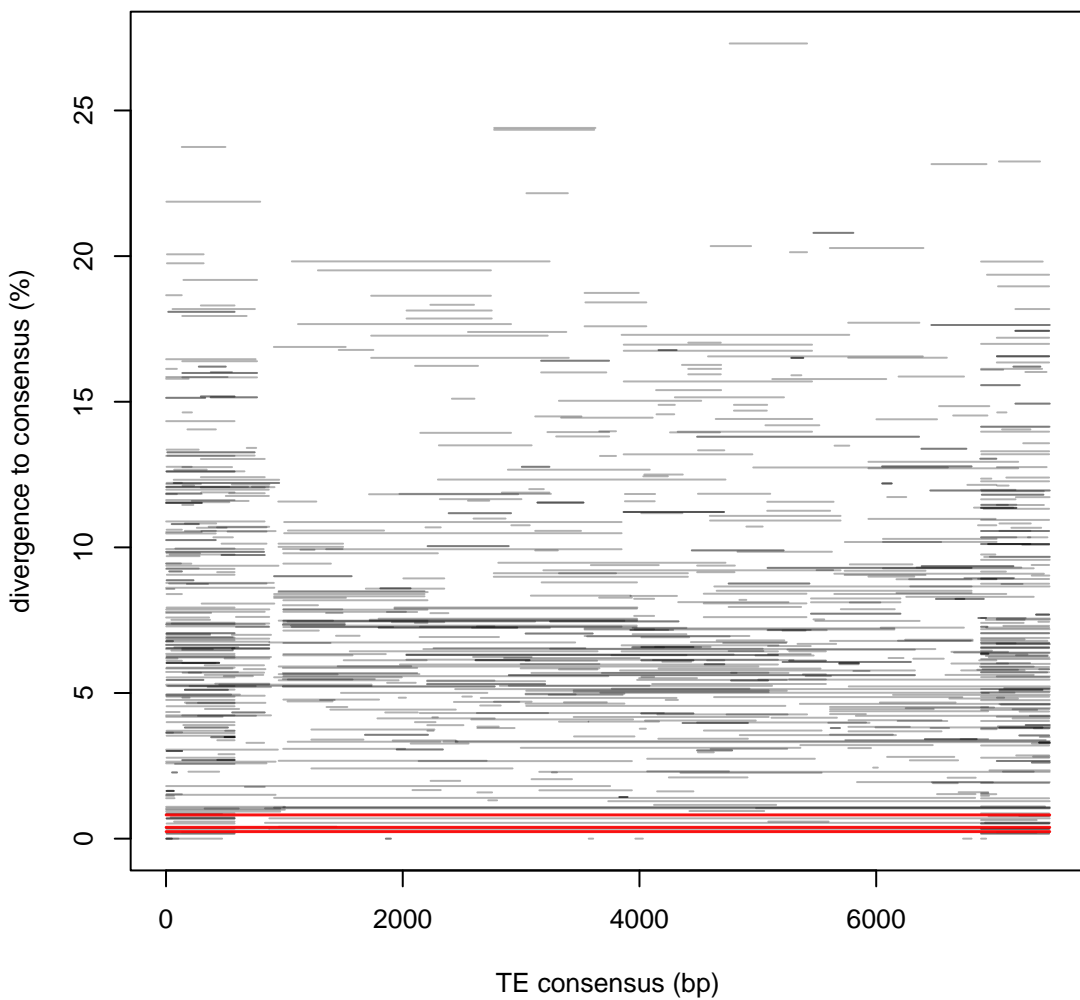
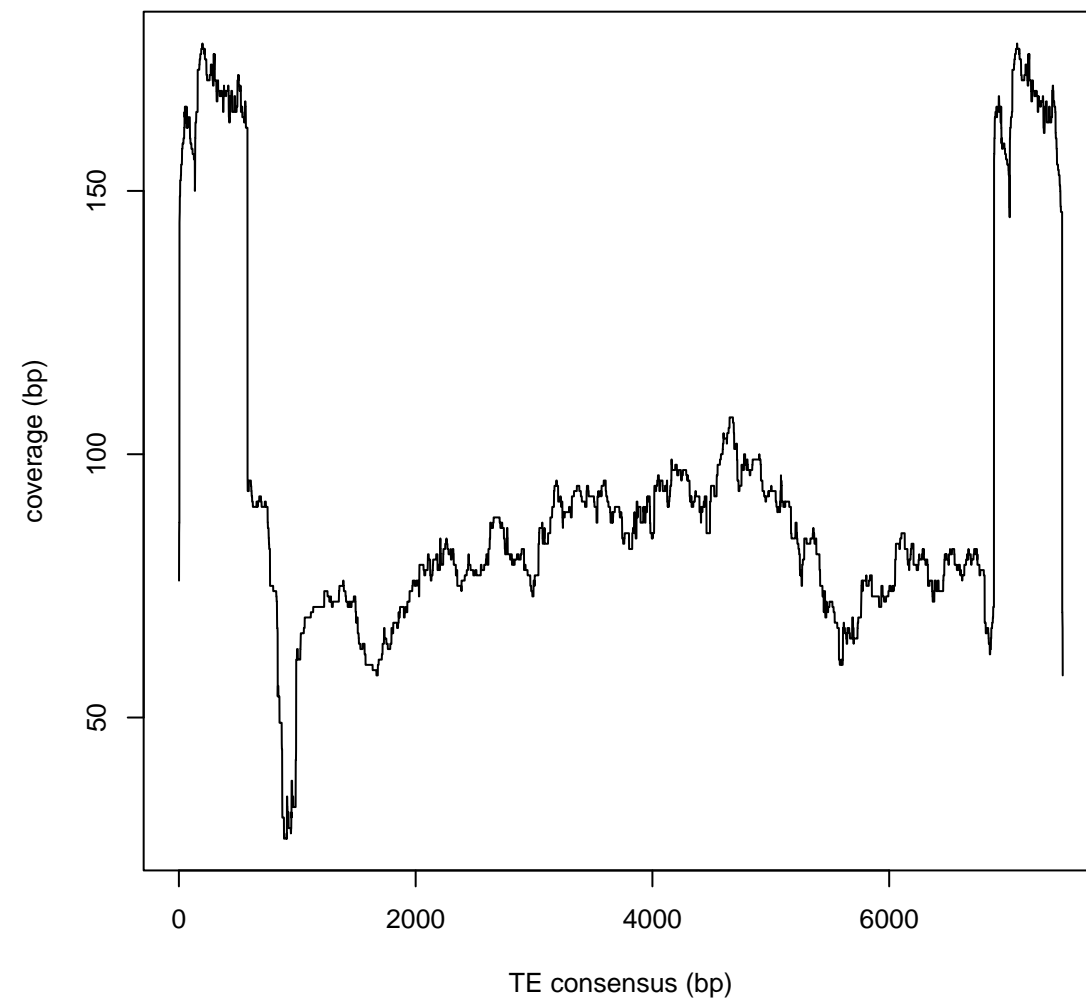


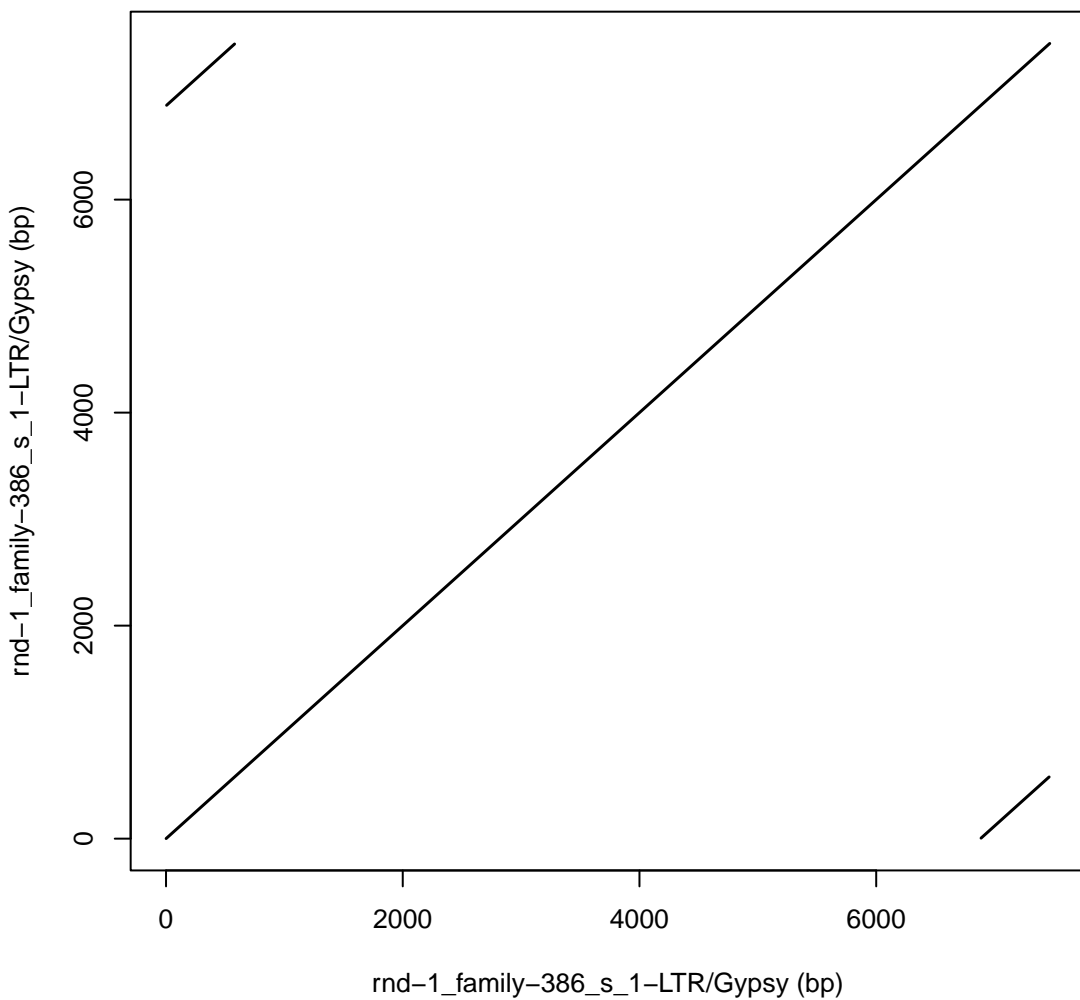
TE: rnd-1_family-386_s_1-LTR/Gypsy
consensus size: 7466bp; fragments: 982; full length: 3 (≥ 6719.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

