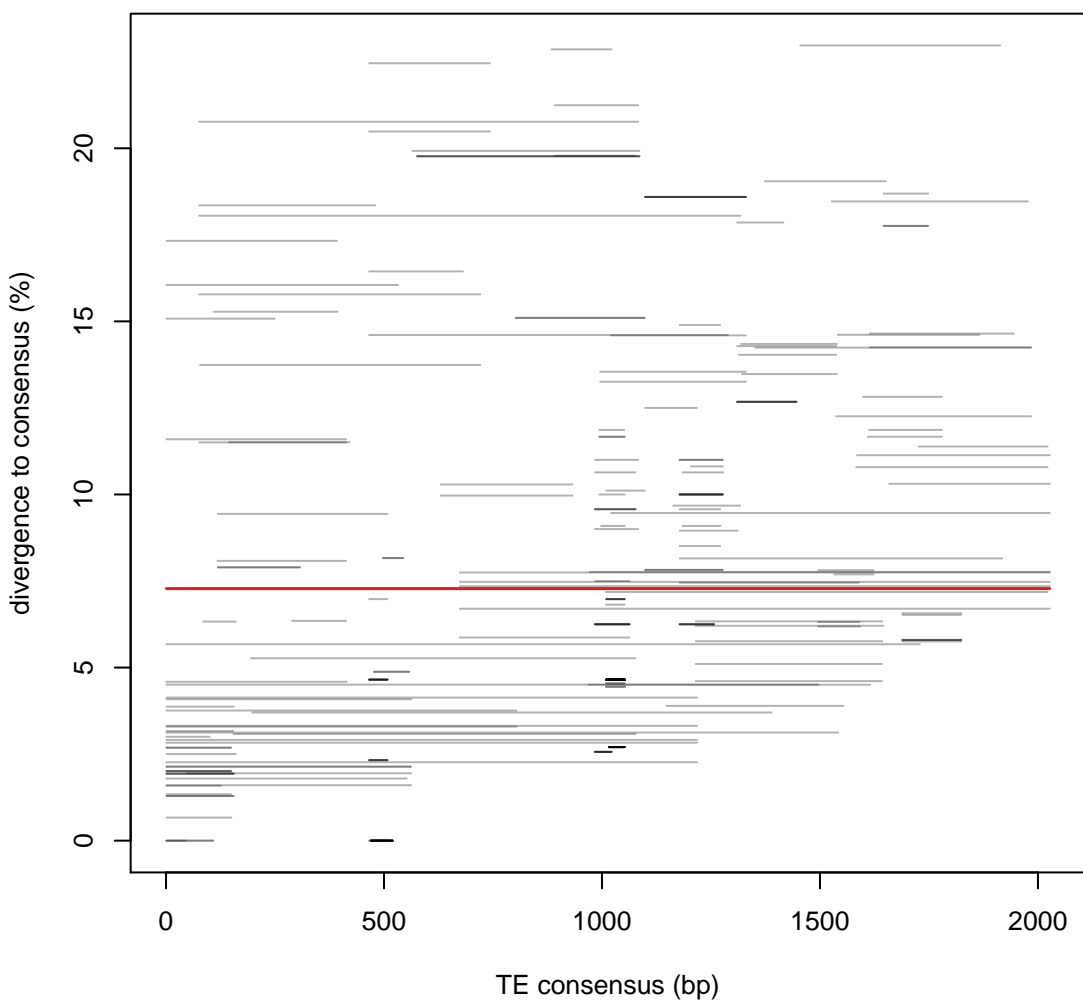
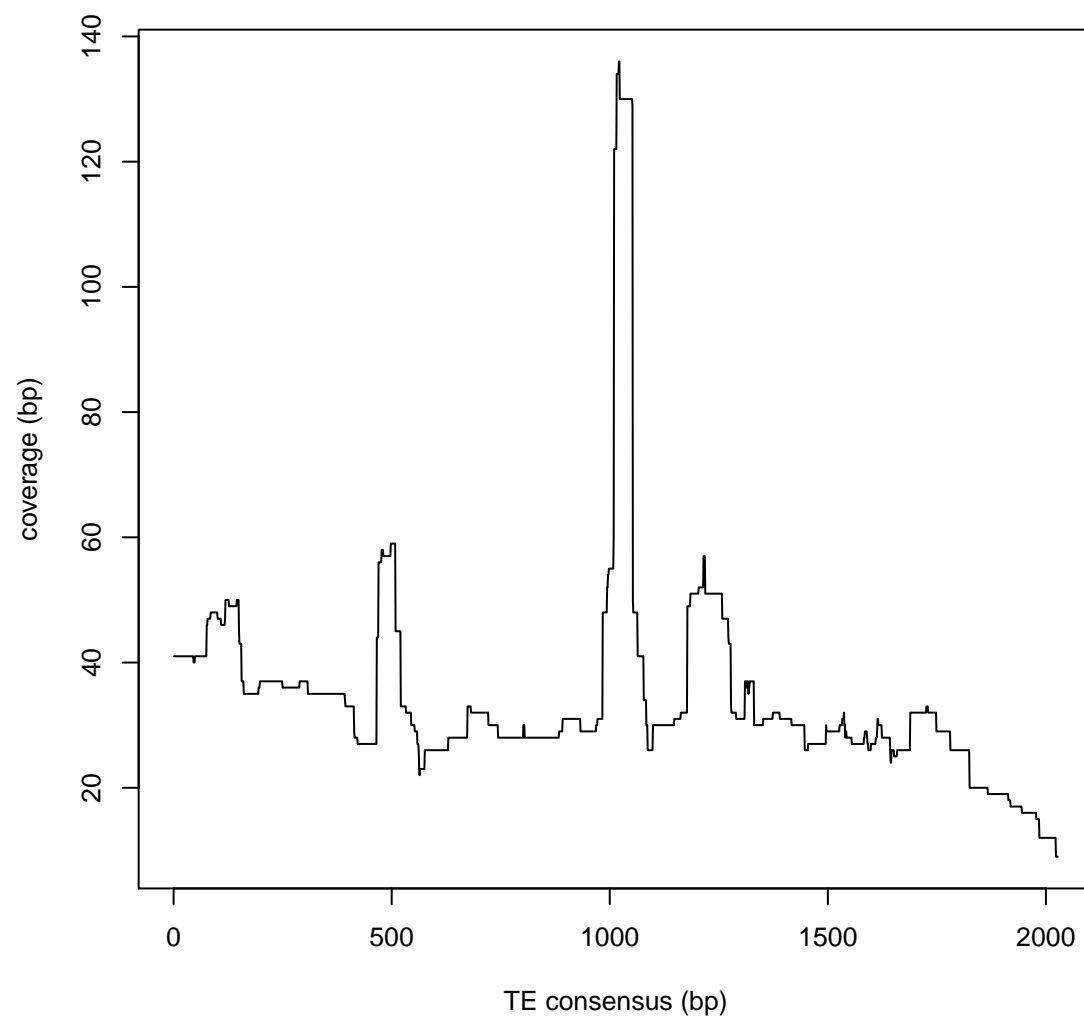


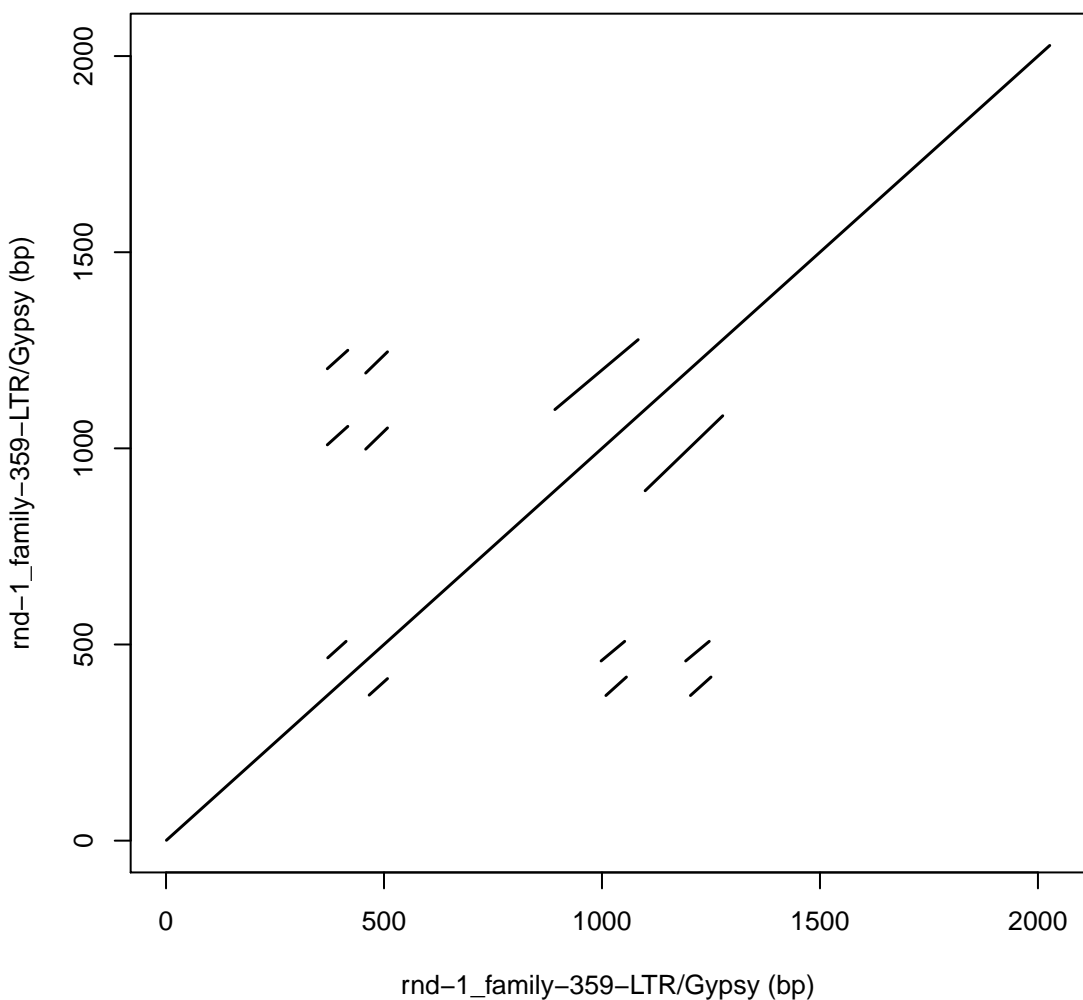
TE: rnd-1_family-359-LTR/Gypsy
consensus size: 2027bp; fragments: 286; full length: 1 (>=1824.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

