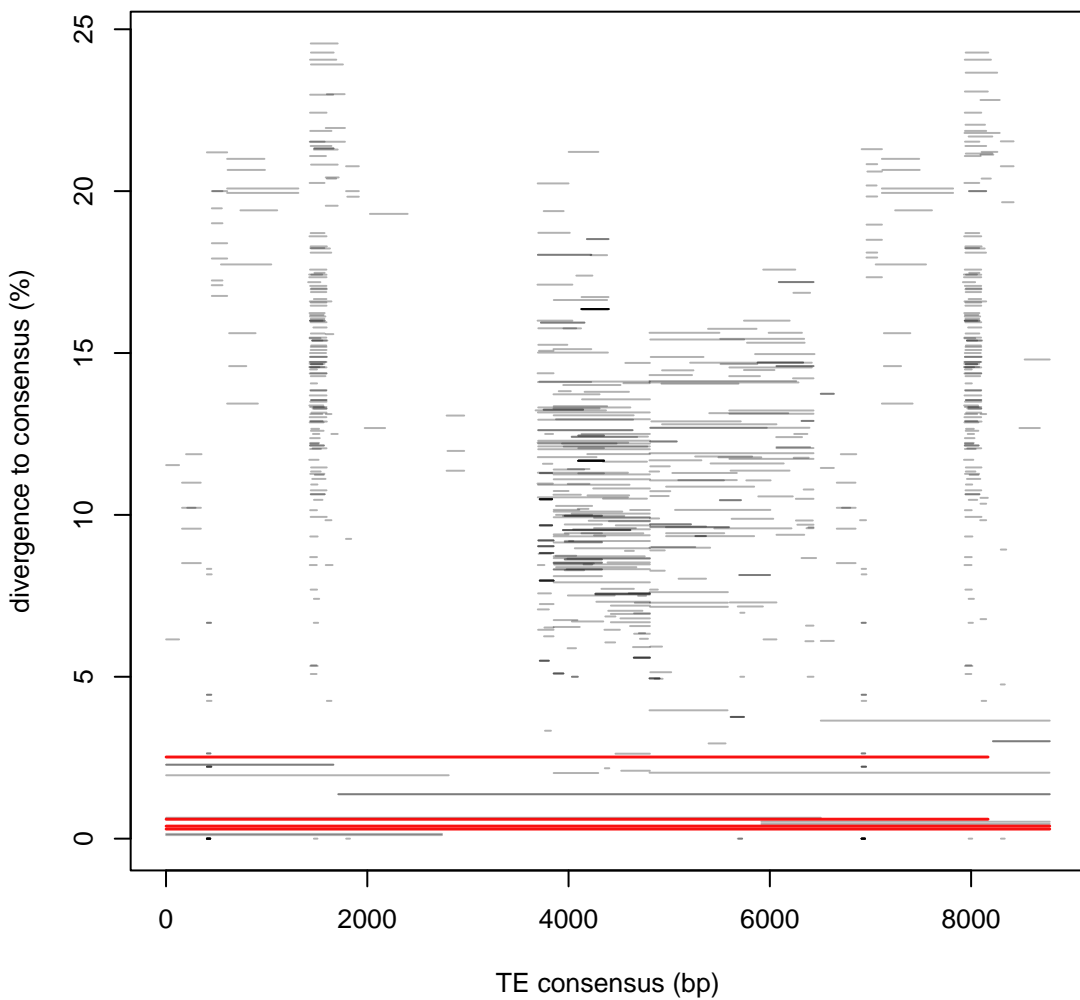
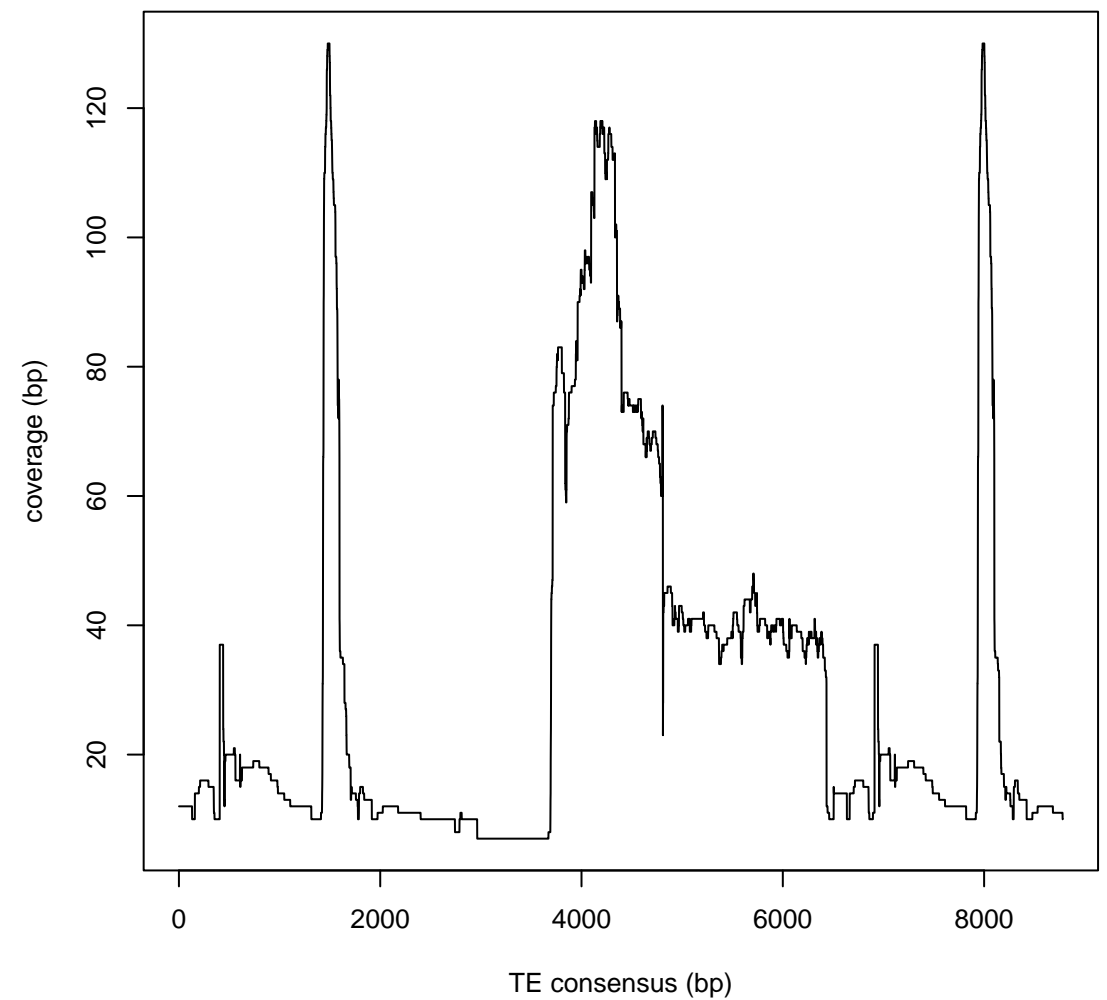


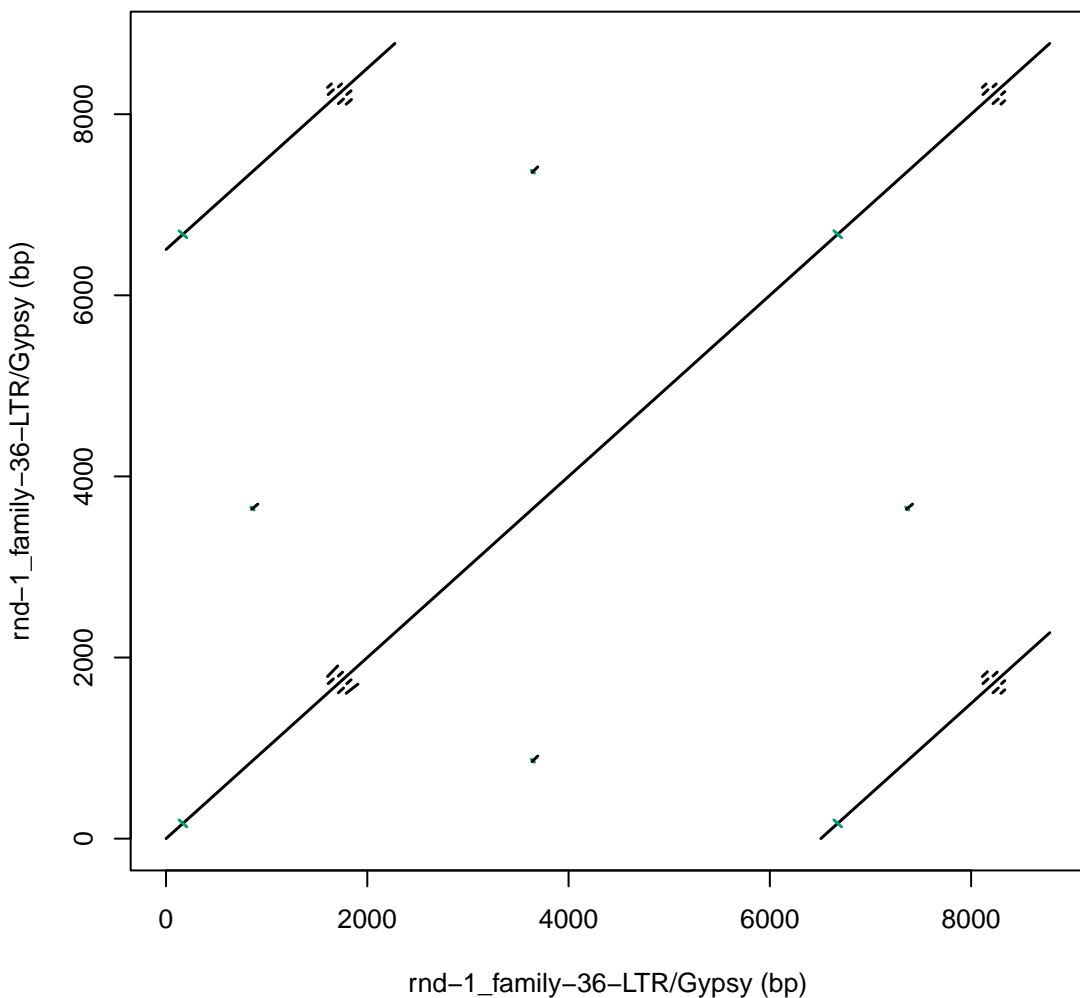
TE: rnd-1_family-36-LTR/Gypsy
consensus size: 8781bp; fragments: 752; full length: 4 (>=7902.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

