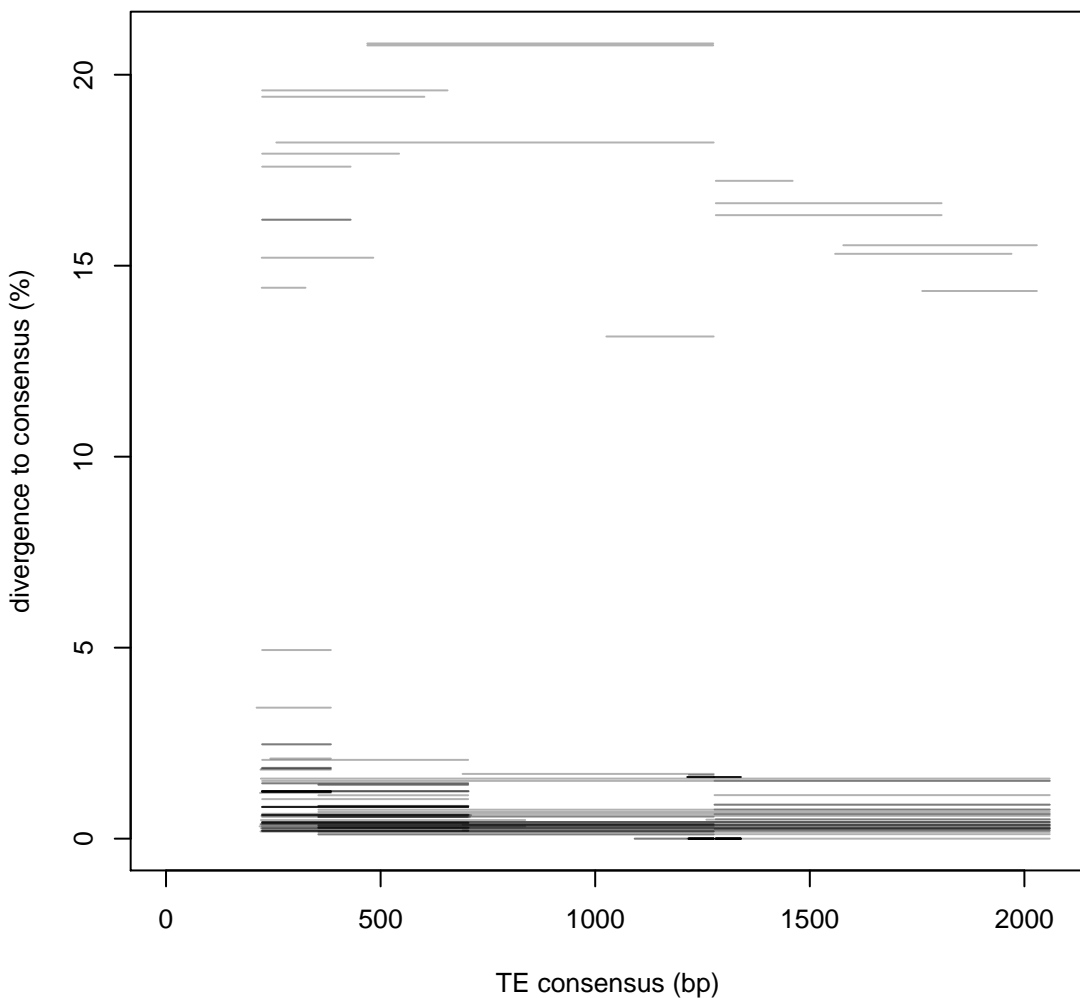
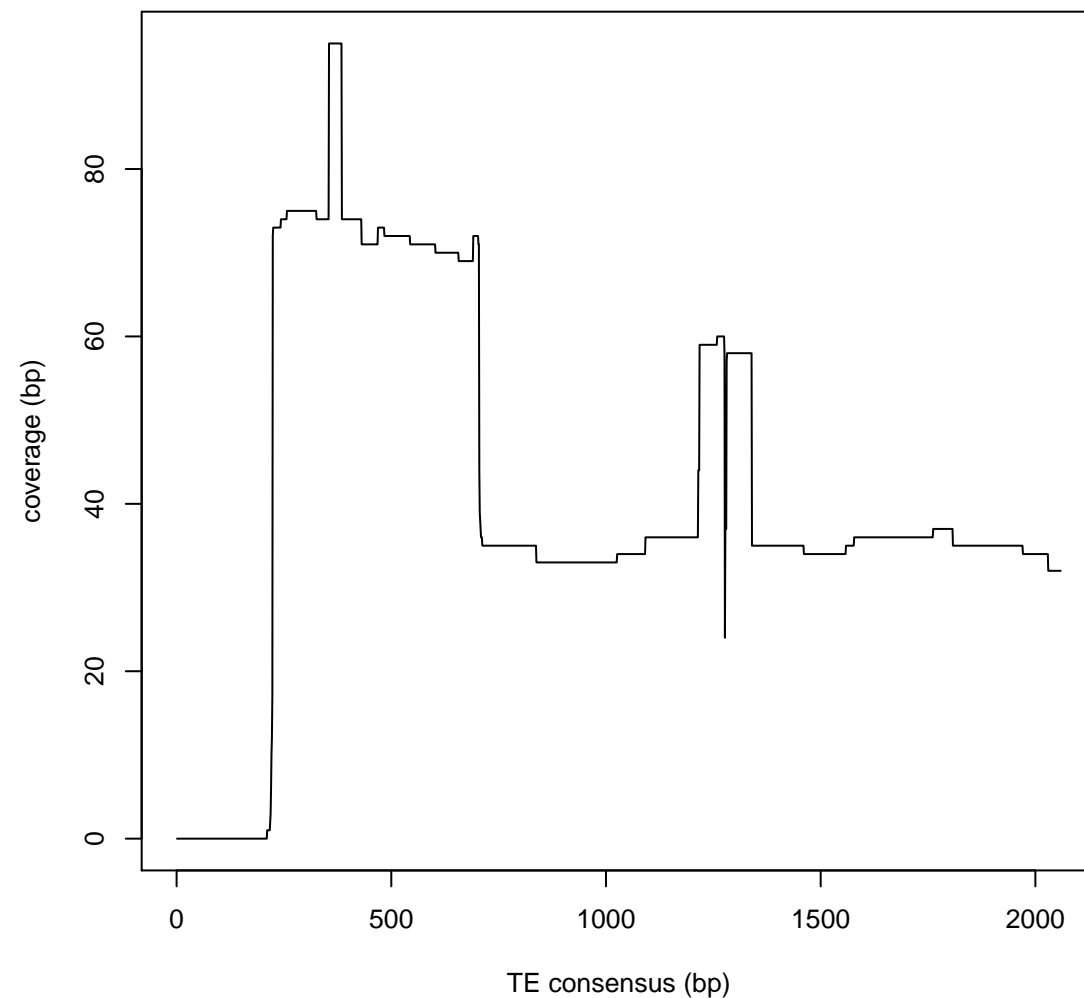


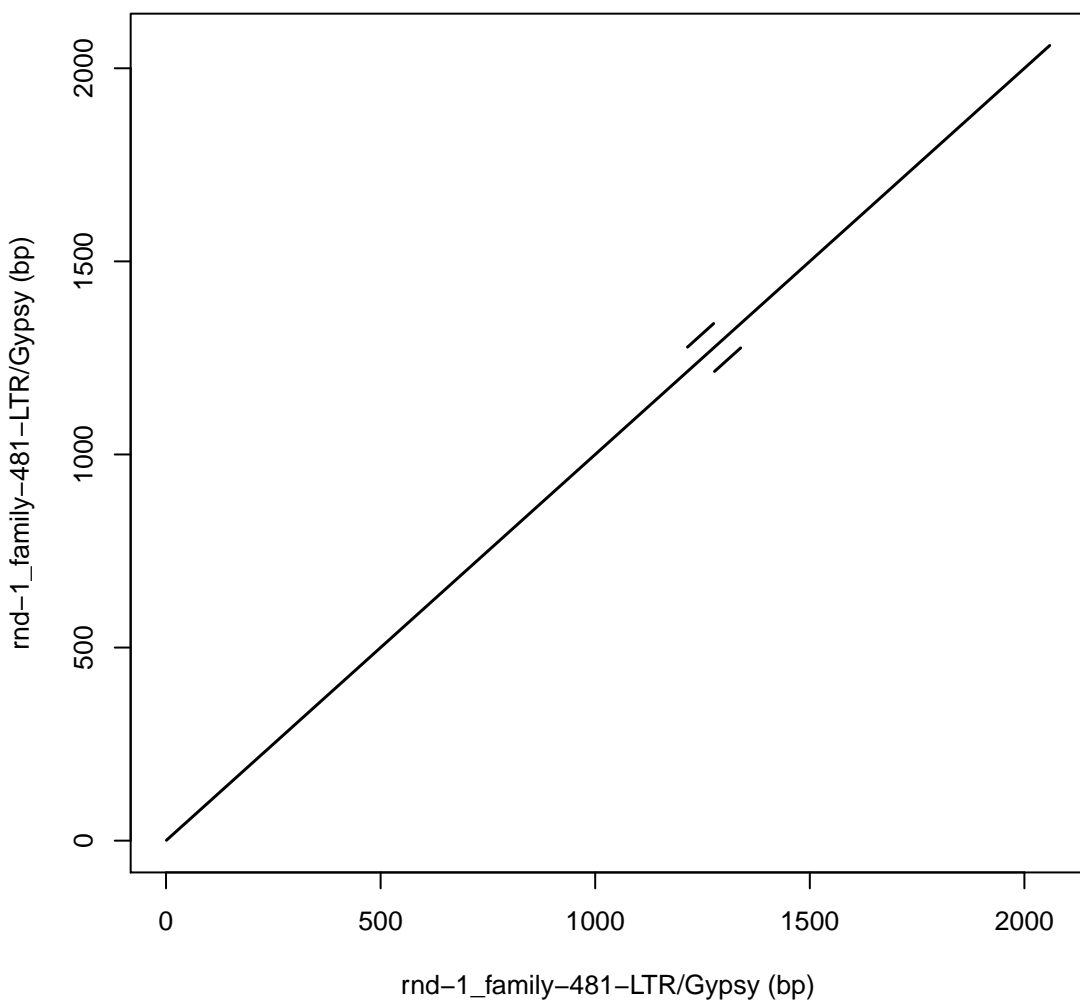
TE: rnd-1_family-481-LTR/Gypsy
consensus size: 2059bp; fragments: 165; full length: 0 (>=1853.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

