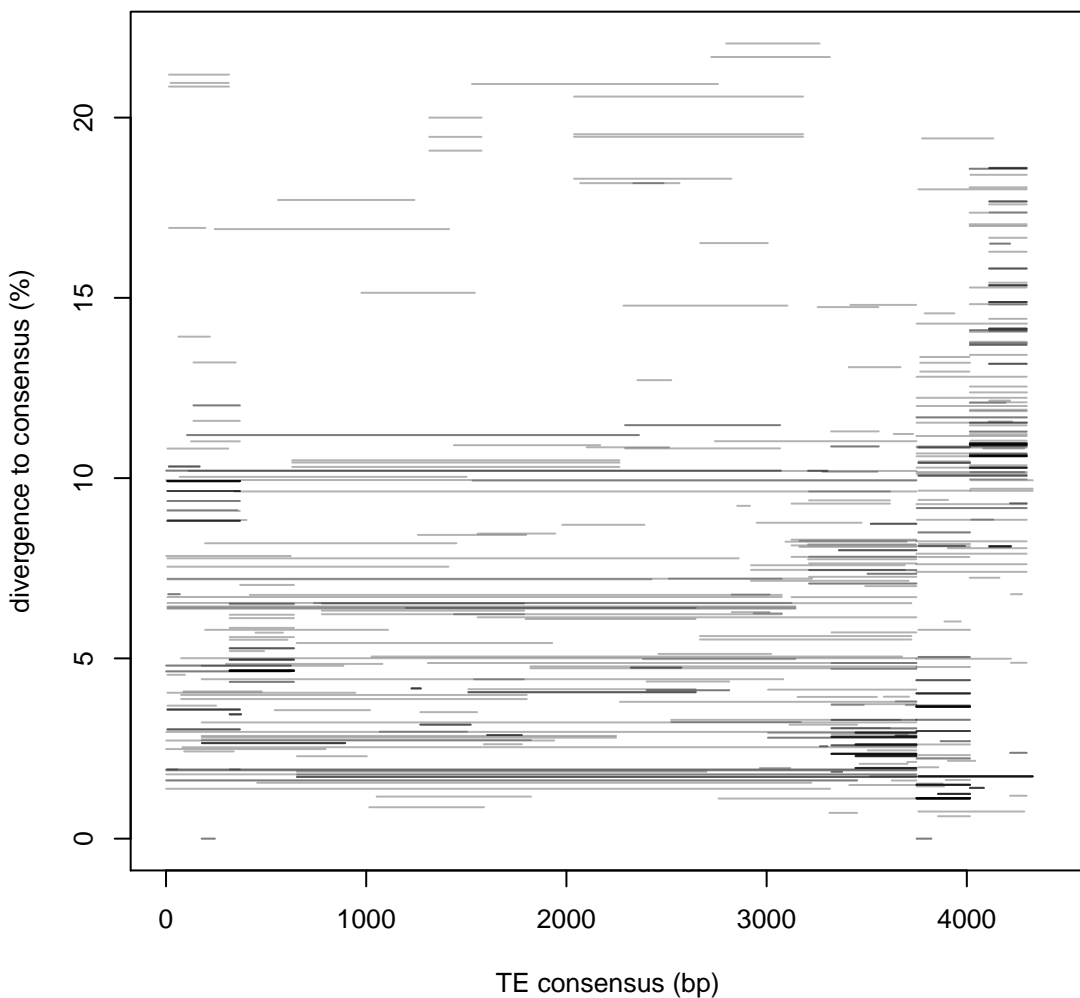
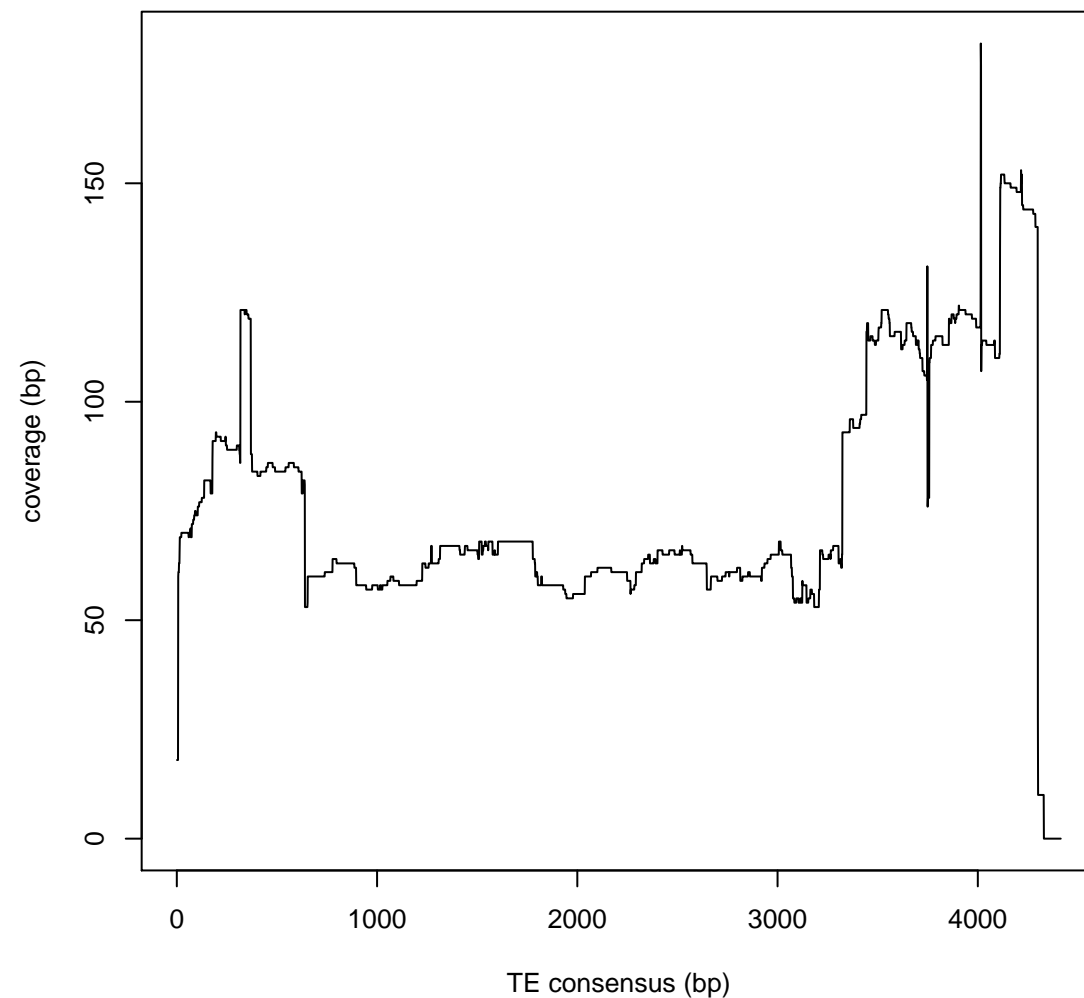


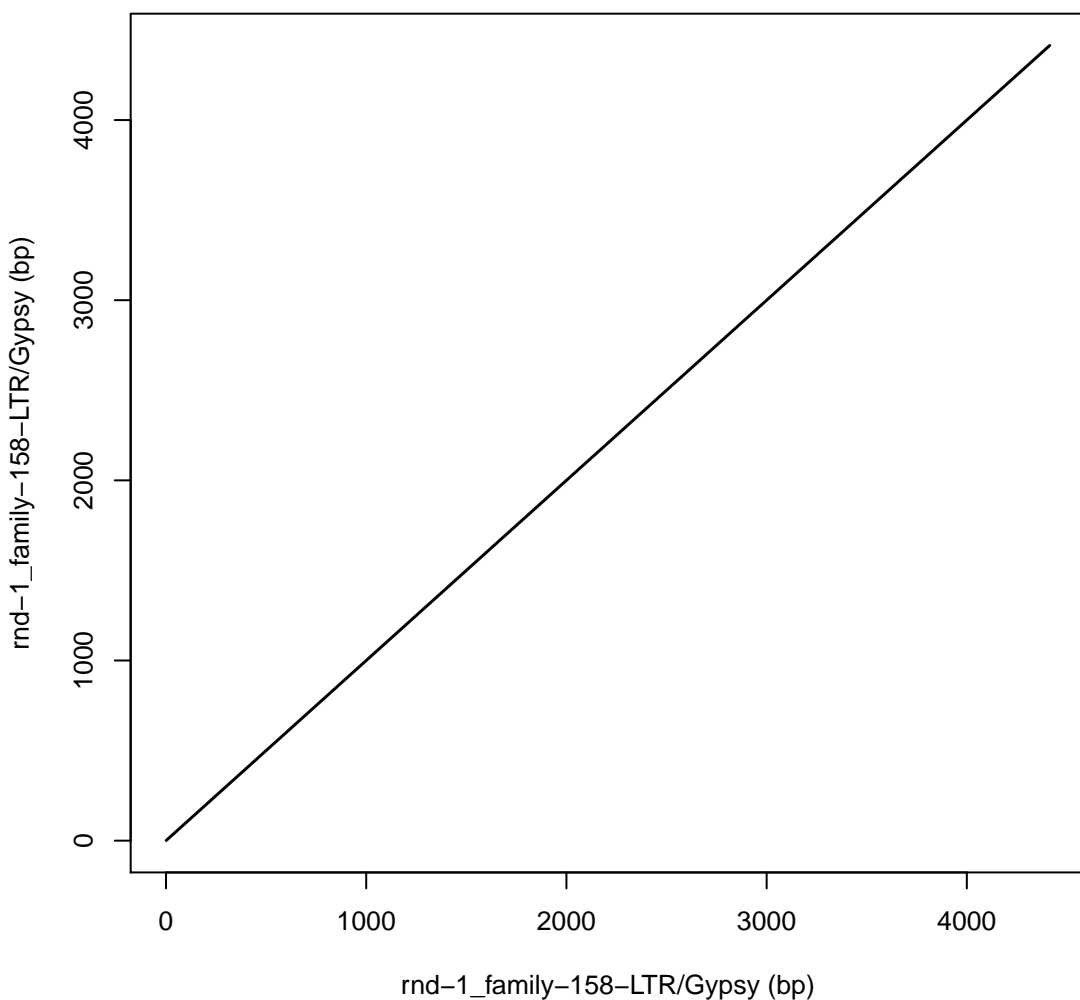
TE: rnd-1_family-158-LTR/Gypsy
consensus size: 4414bp; fragments: 595; full length: 0 (>=3972.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

