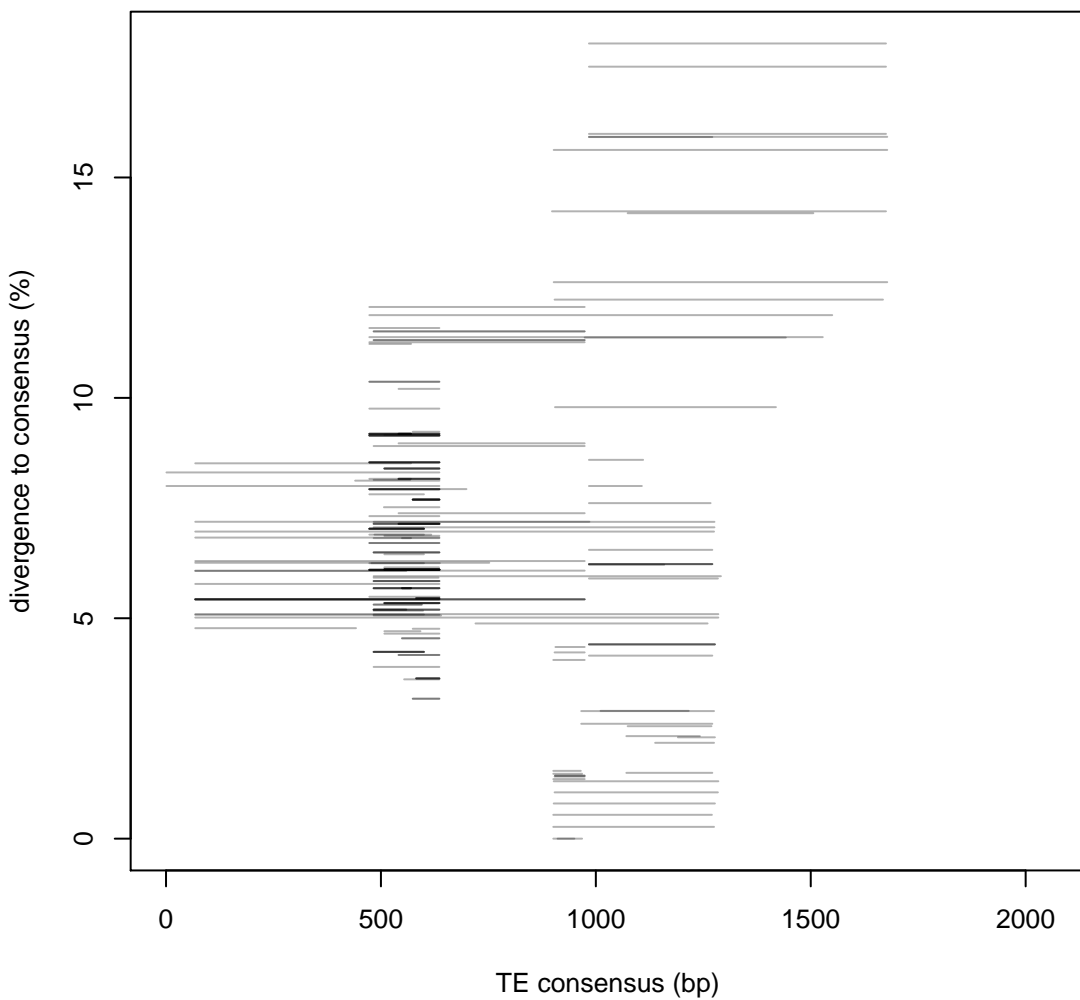
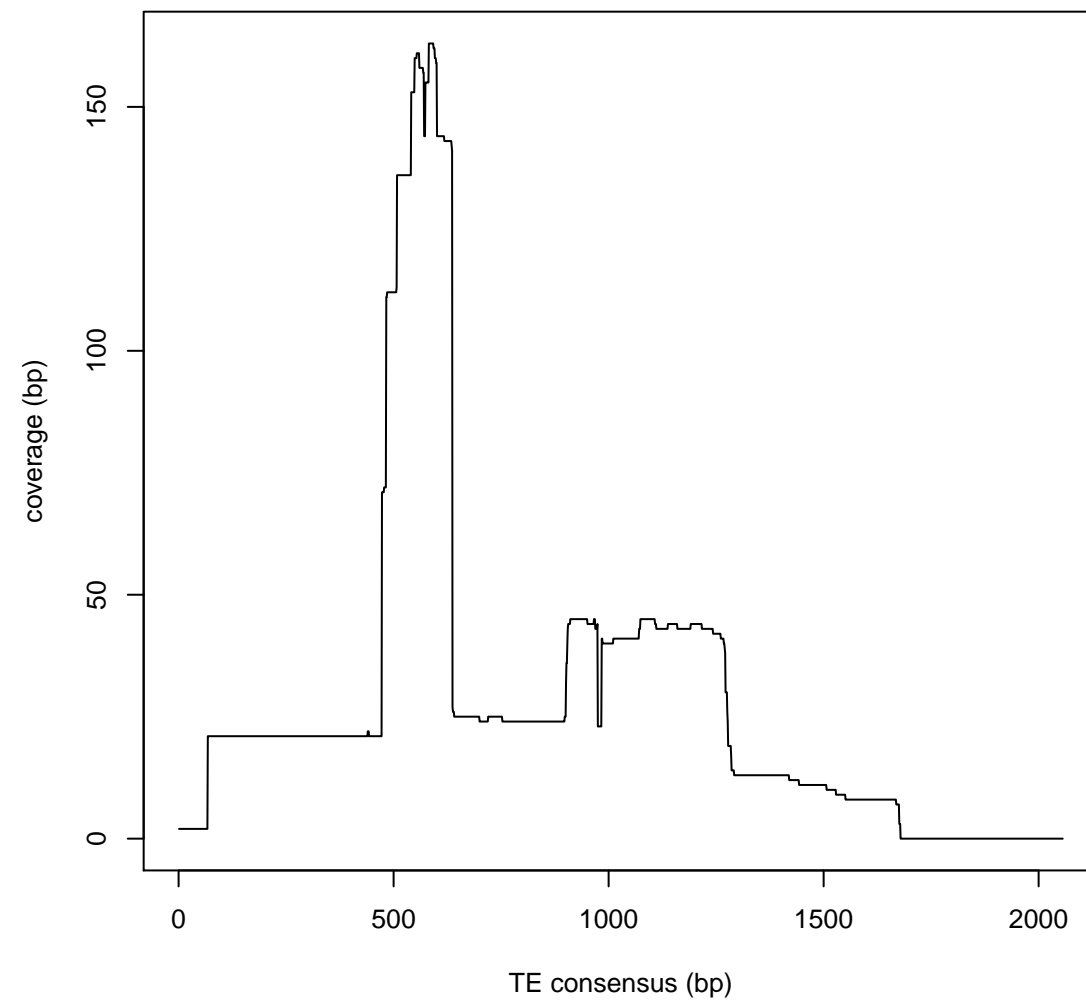


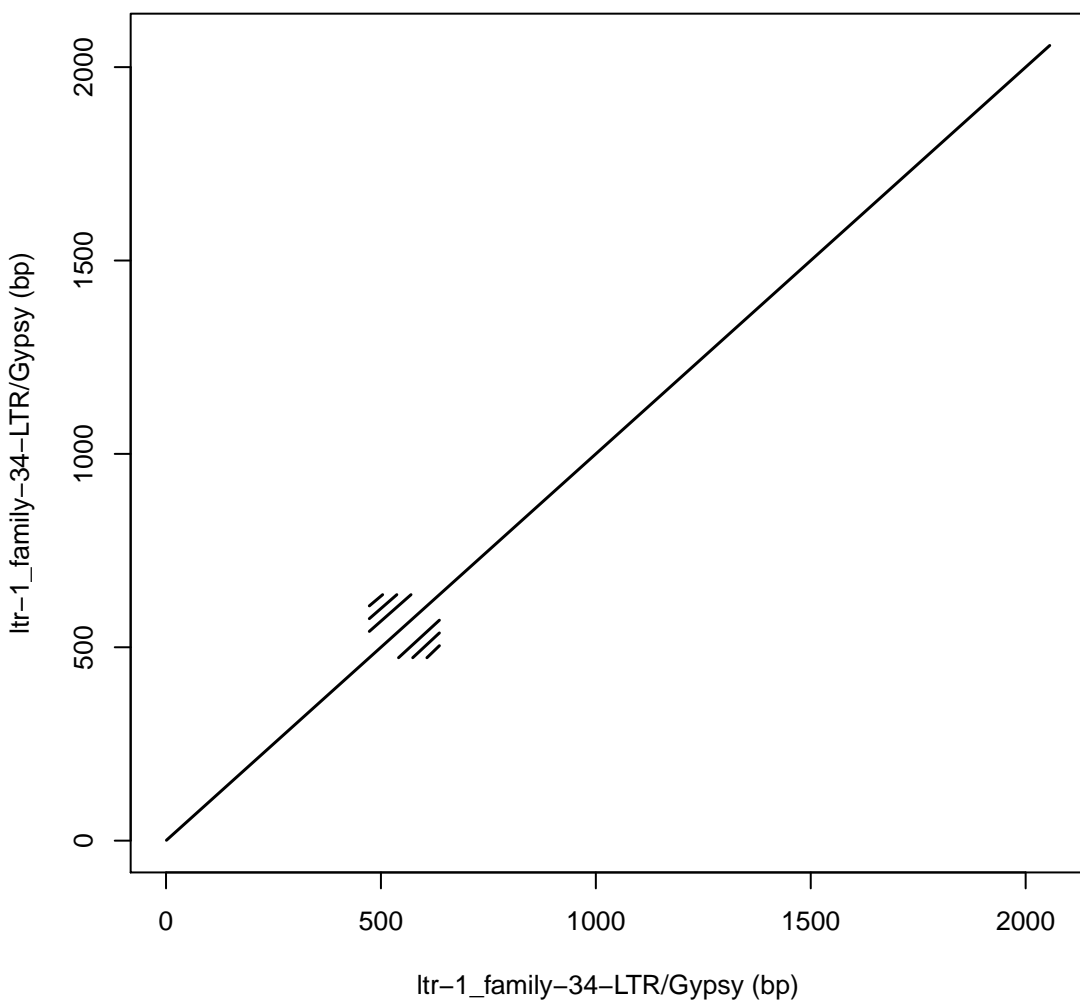
TE: ltr-1_family-34-LTR/Gypsy
consensus size: 2056bp; fragments: 231; full length: 0 (>=1850.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

