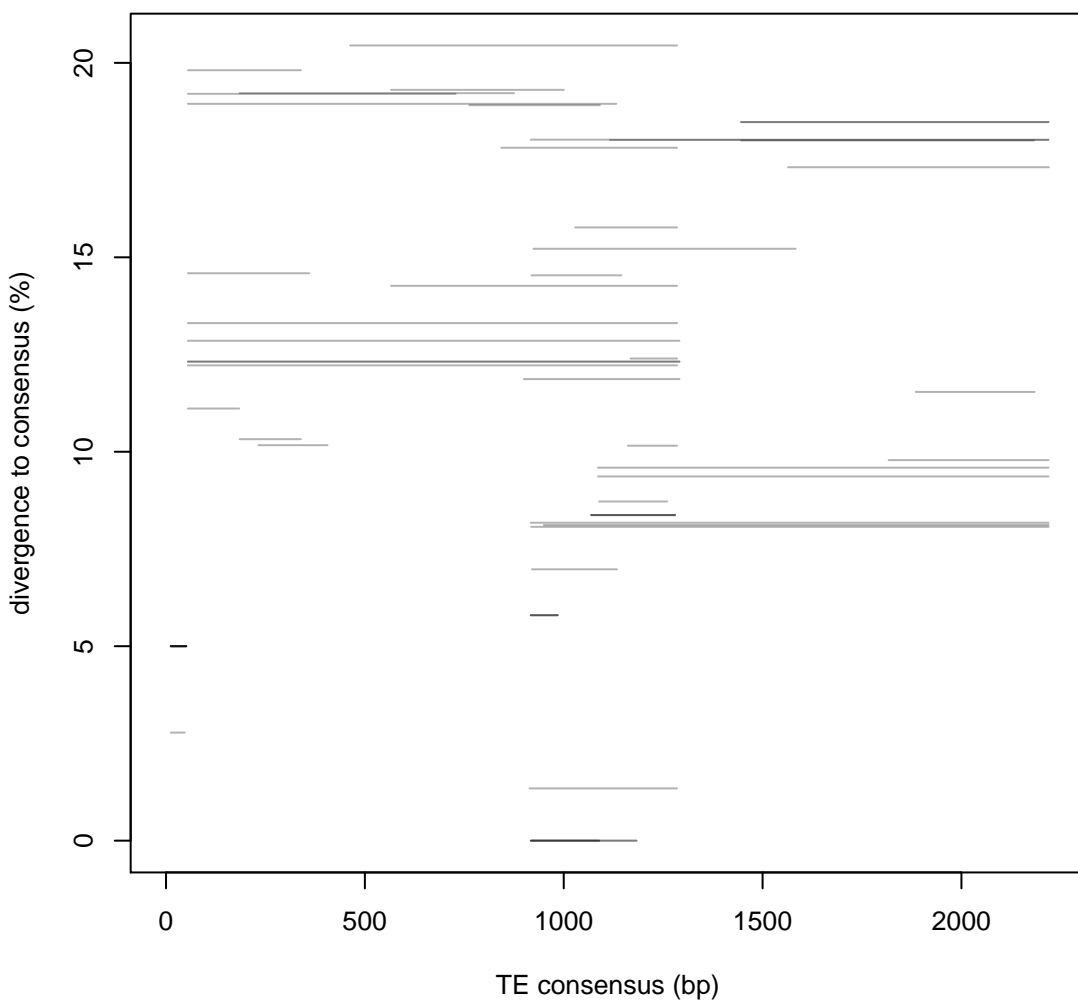
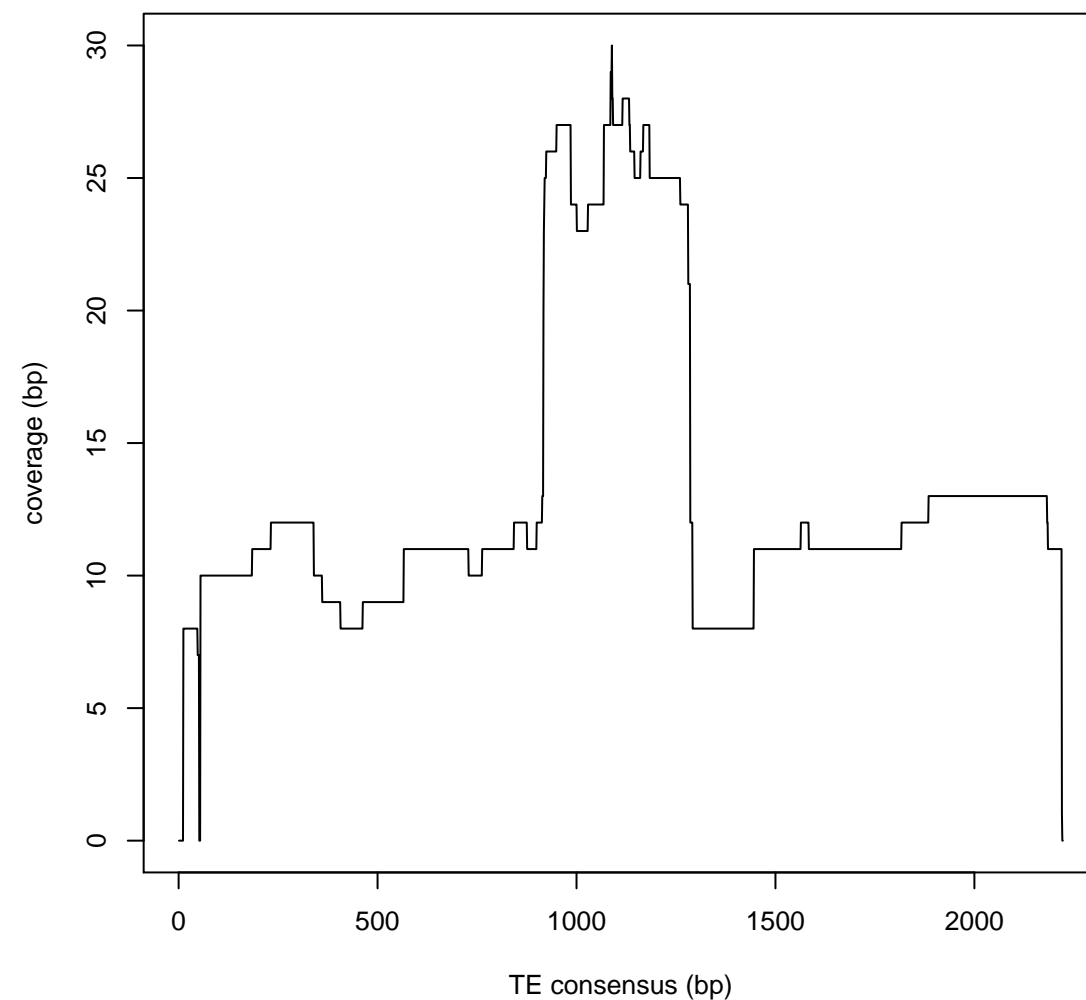


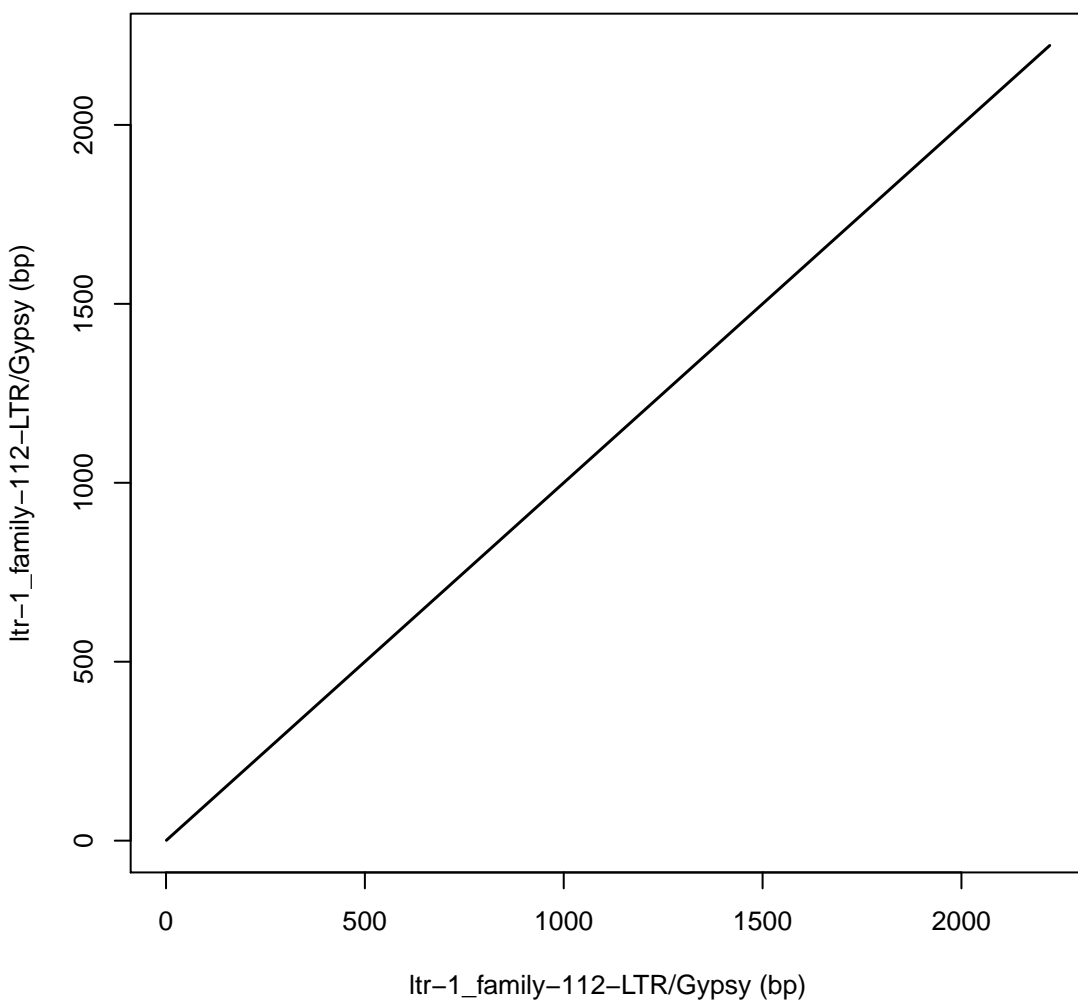
TE: ltr-1_family-112-LTR/Gypsy
consensus size: 2222bp; fragments: 58; full length: 0 (≥ 1999.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

