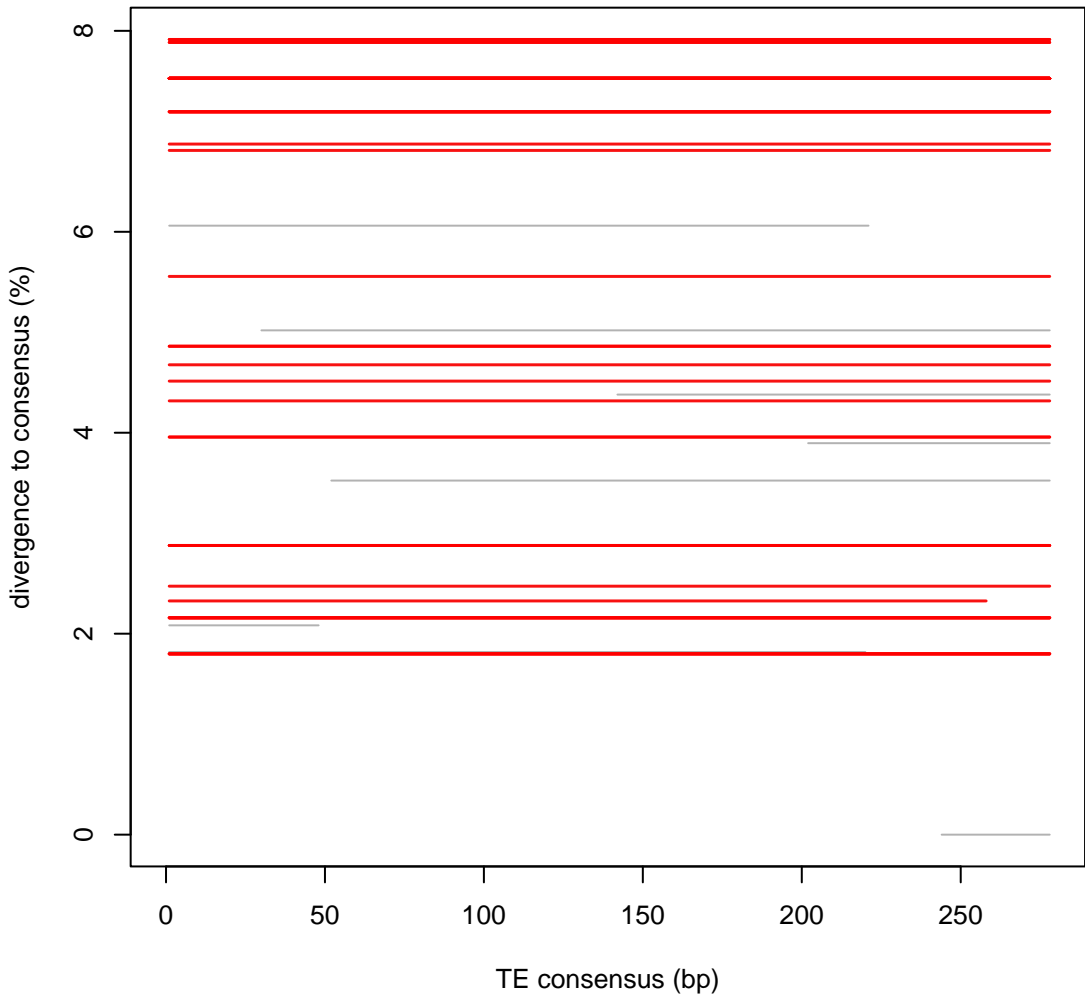
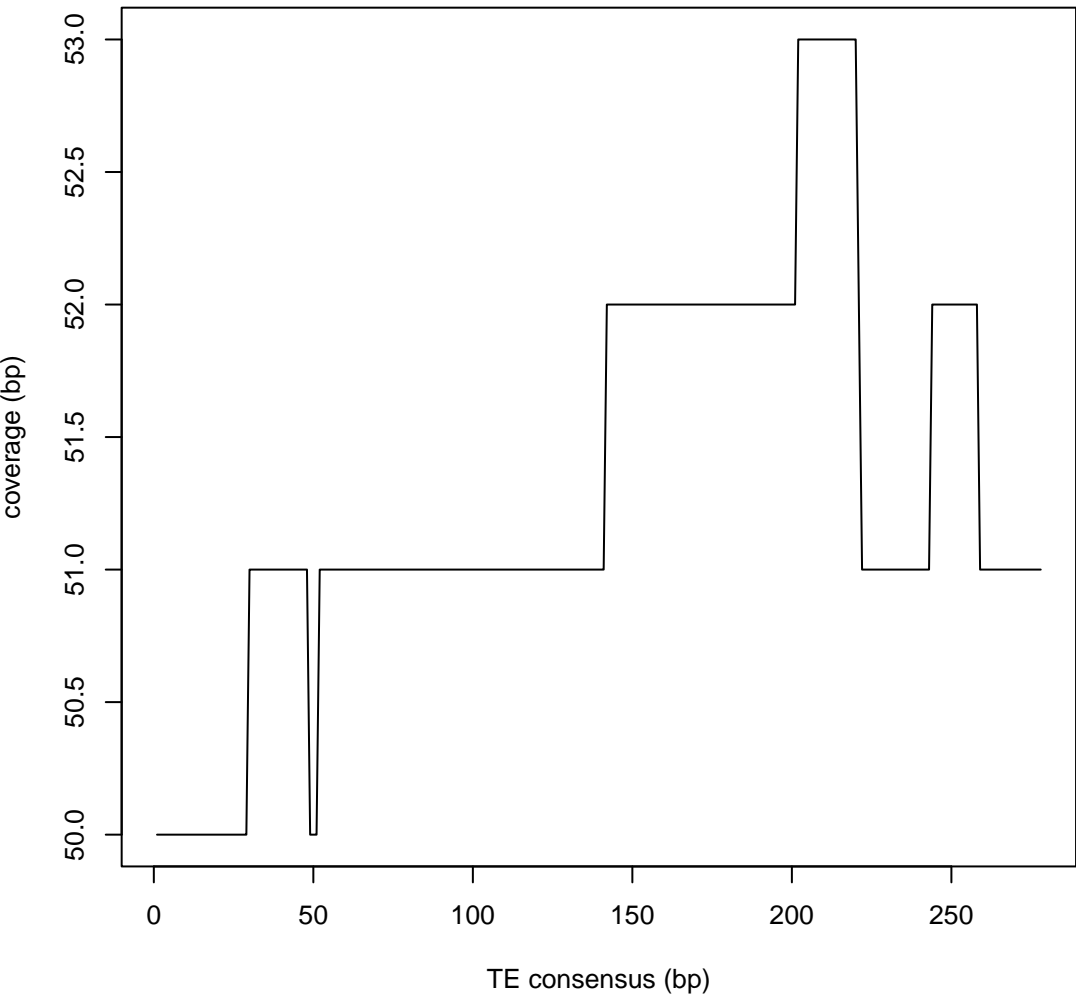


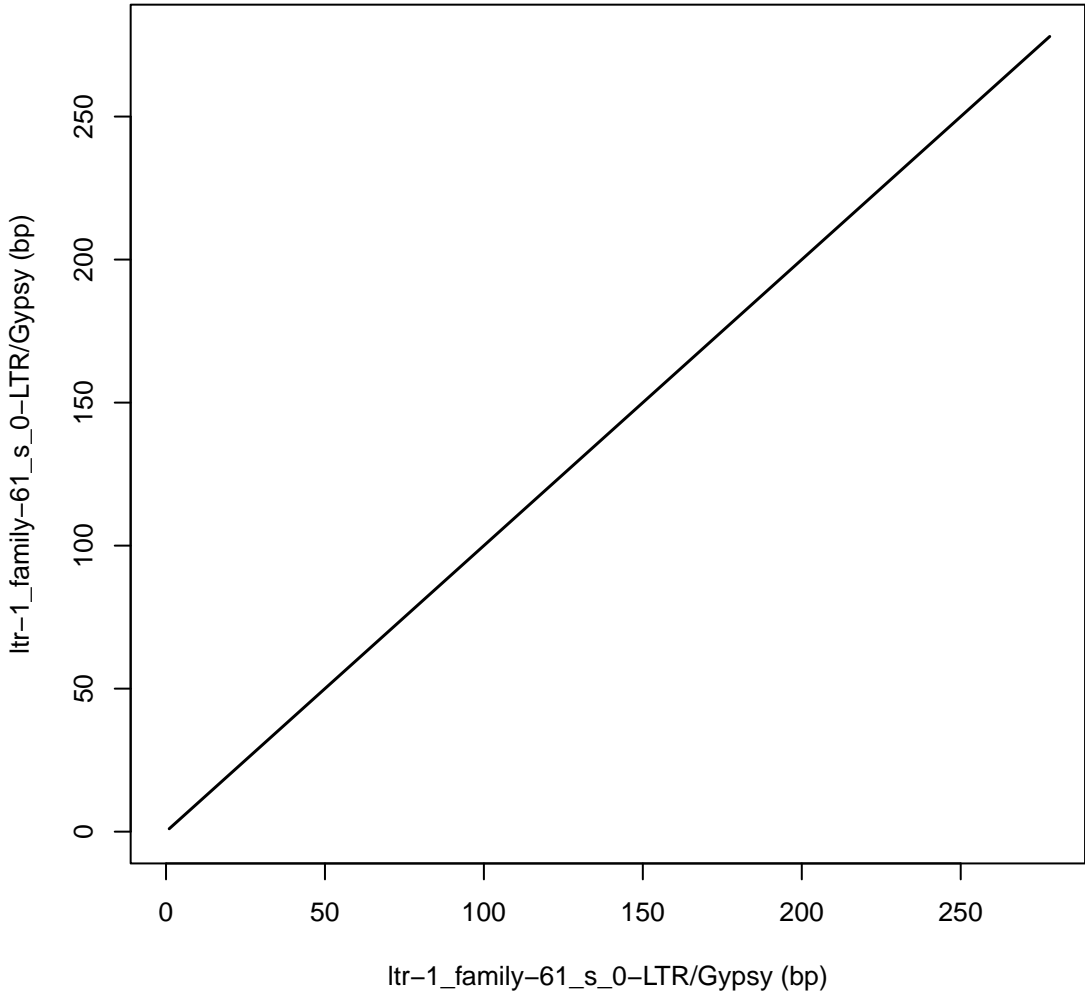
TE: ltr-1_family-61_s_0-LTR/Gypsy
consensus size: 278bp; fragments: 55; full length: 47 (>=250.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

