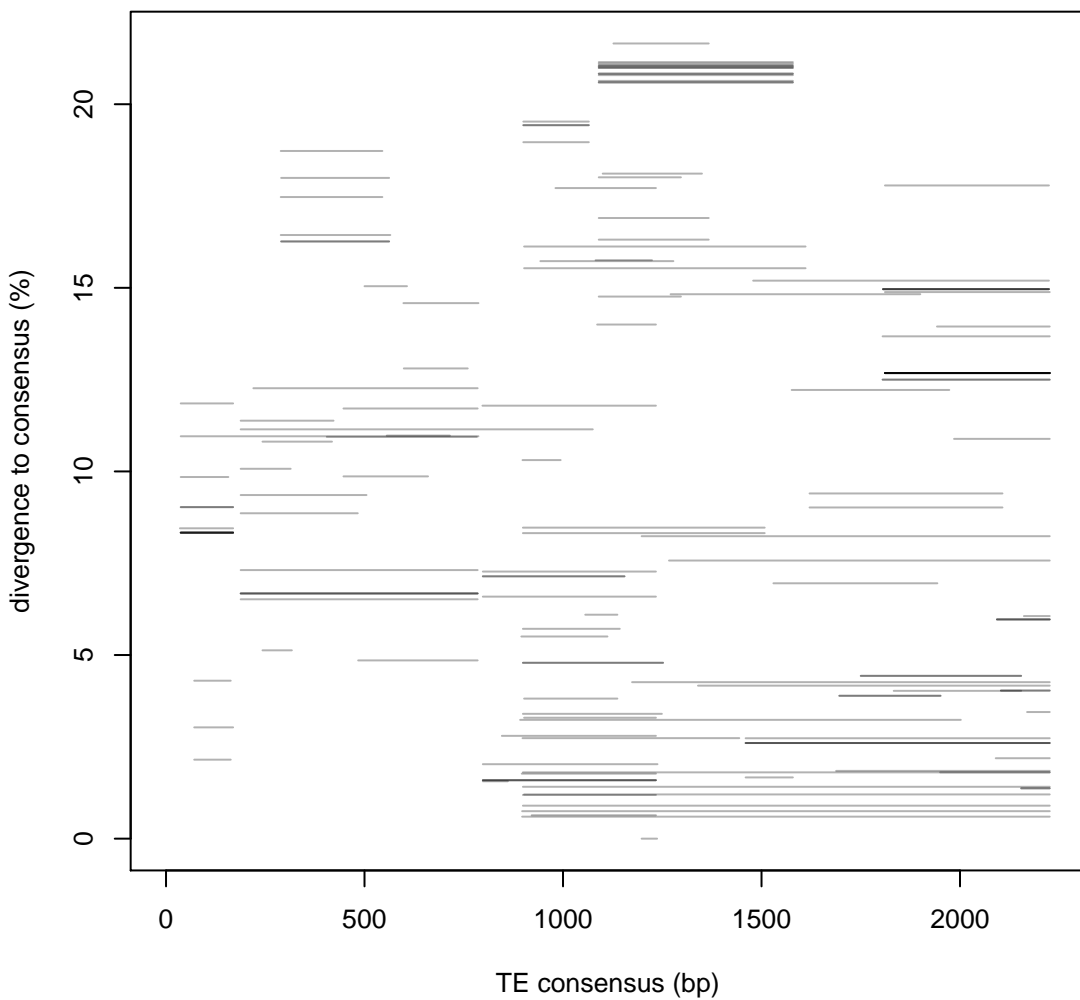
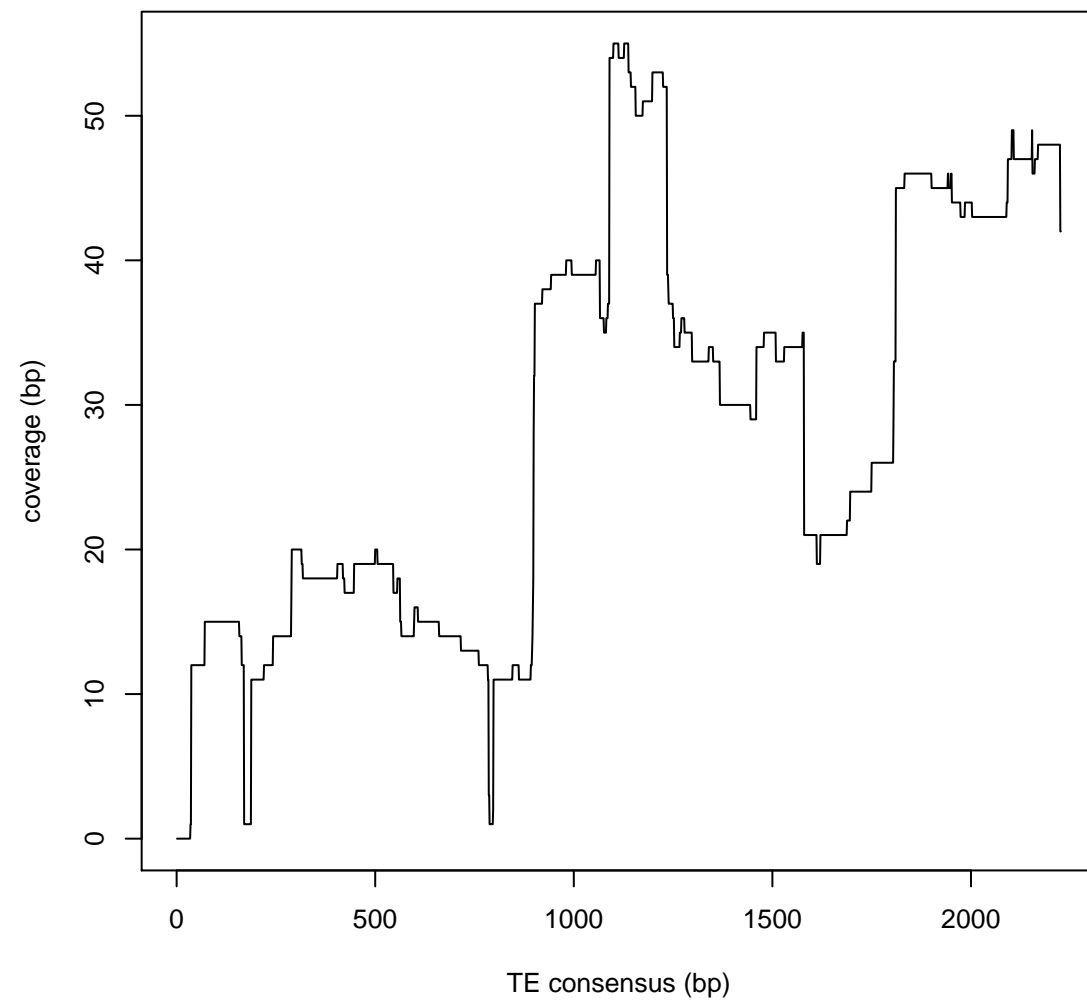


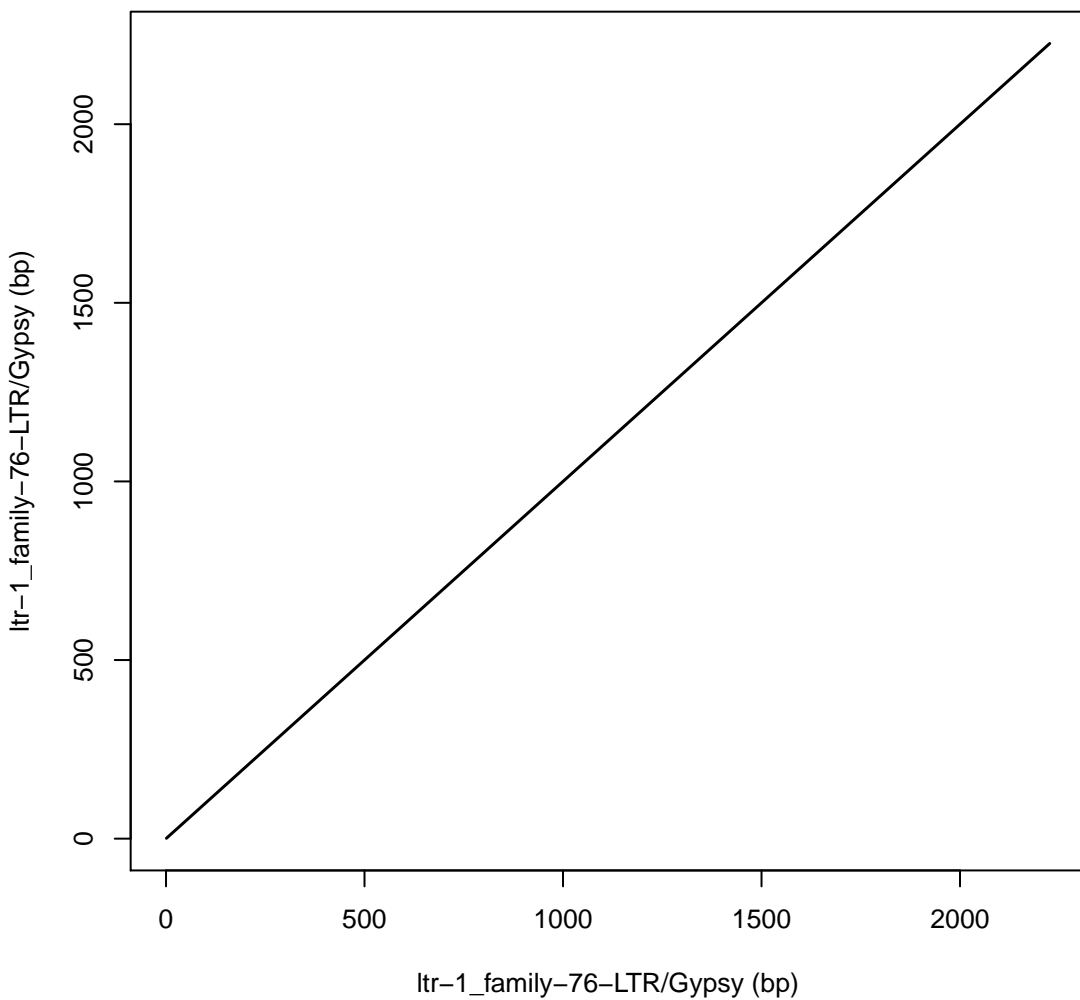
TE: ltr-1_family-76-LTR/Gypsy
consensus size: 2226bp; fragments: 158; full length: 0 (>=2003.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

