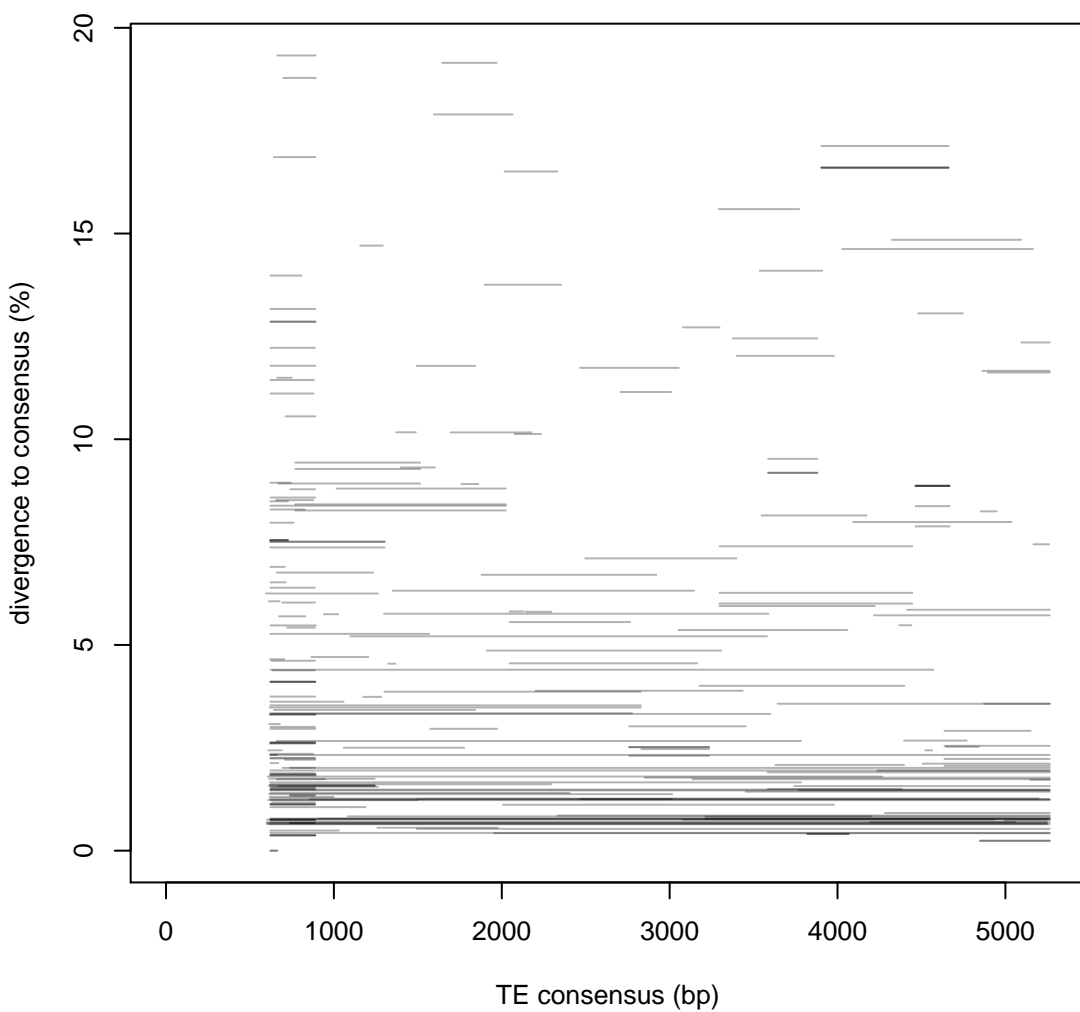
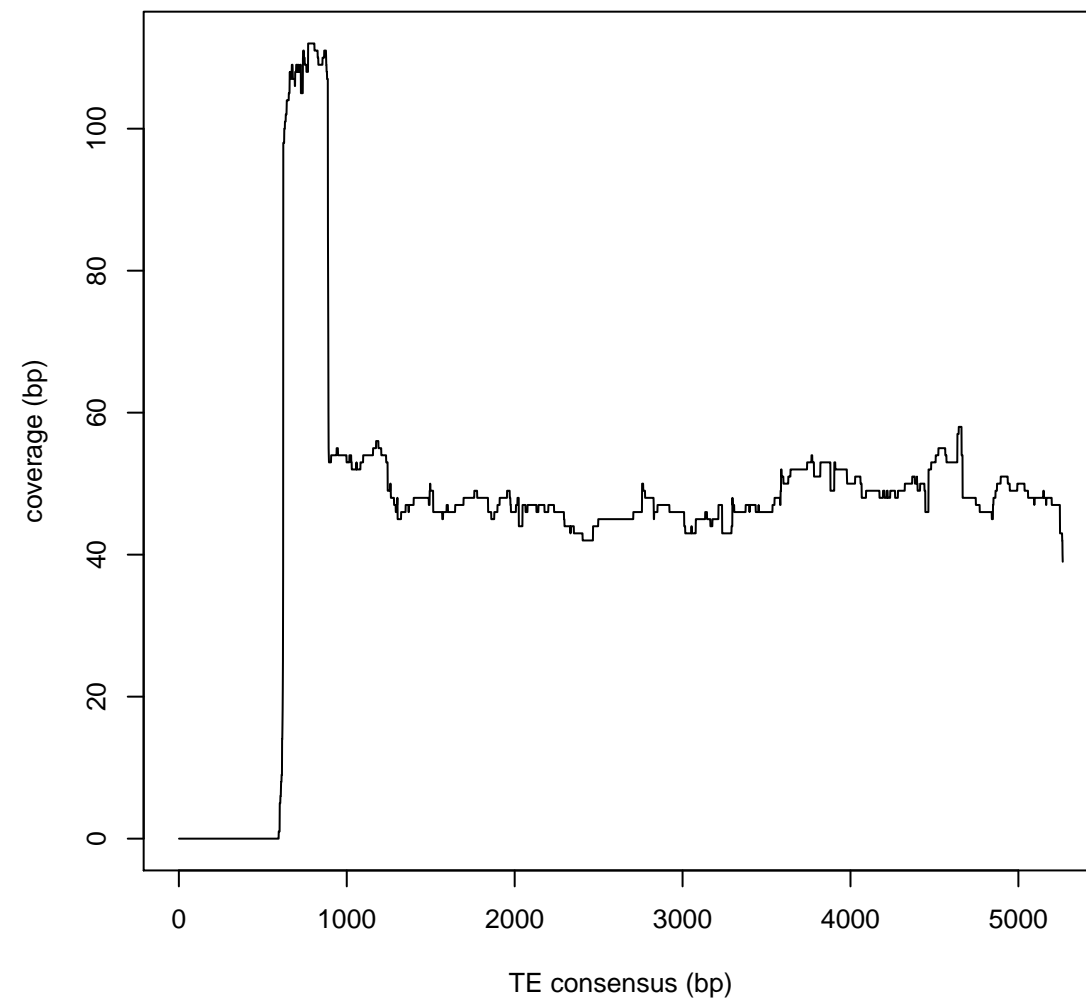


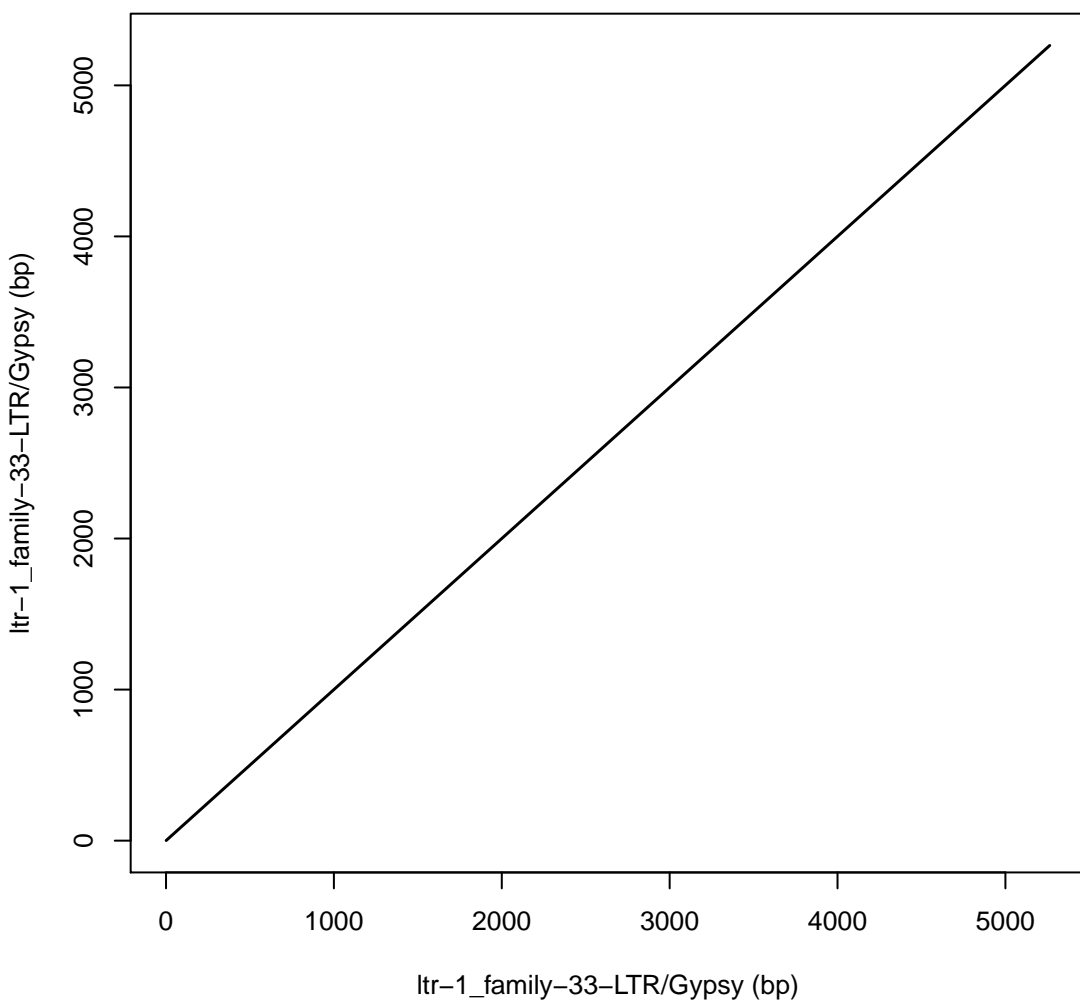
TE: ltr-1_family-33-LTR/Gypsy
consensus size: 5264bp; fragments: 243; full length: 0 (>=4737.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

