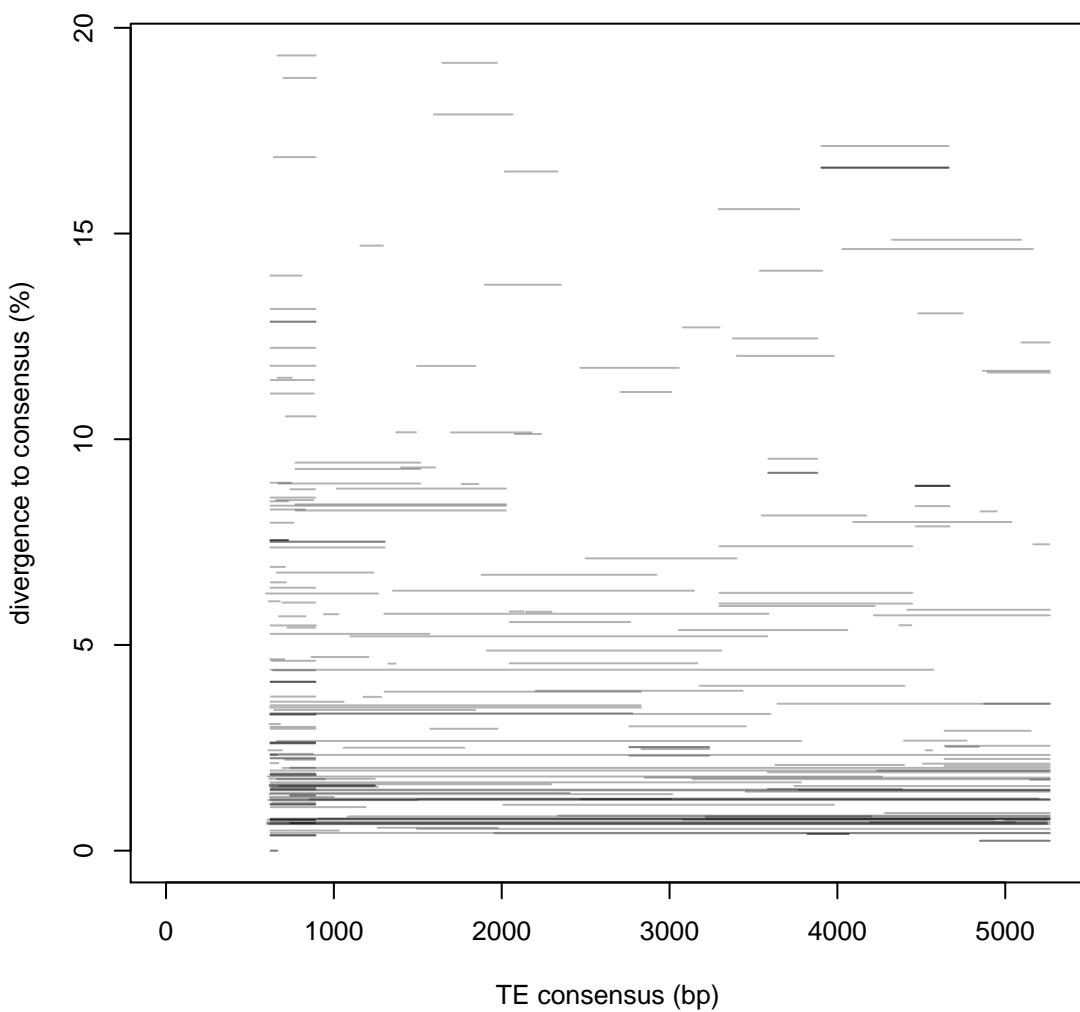
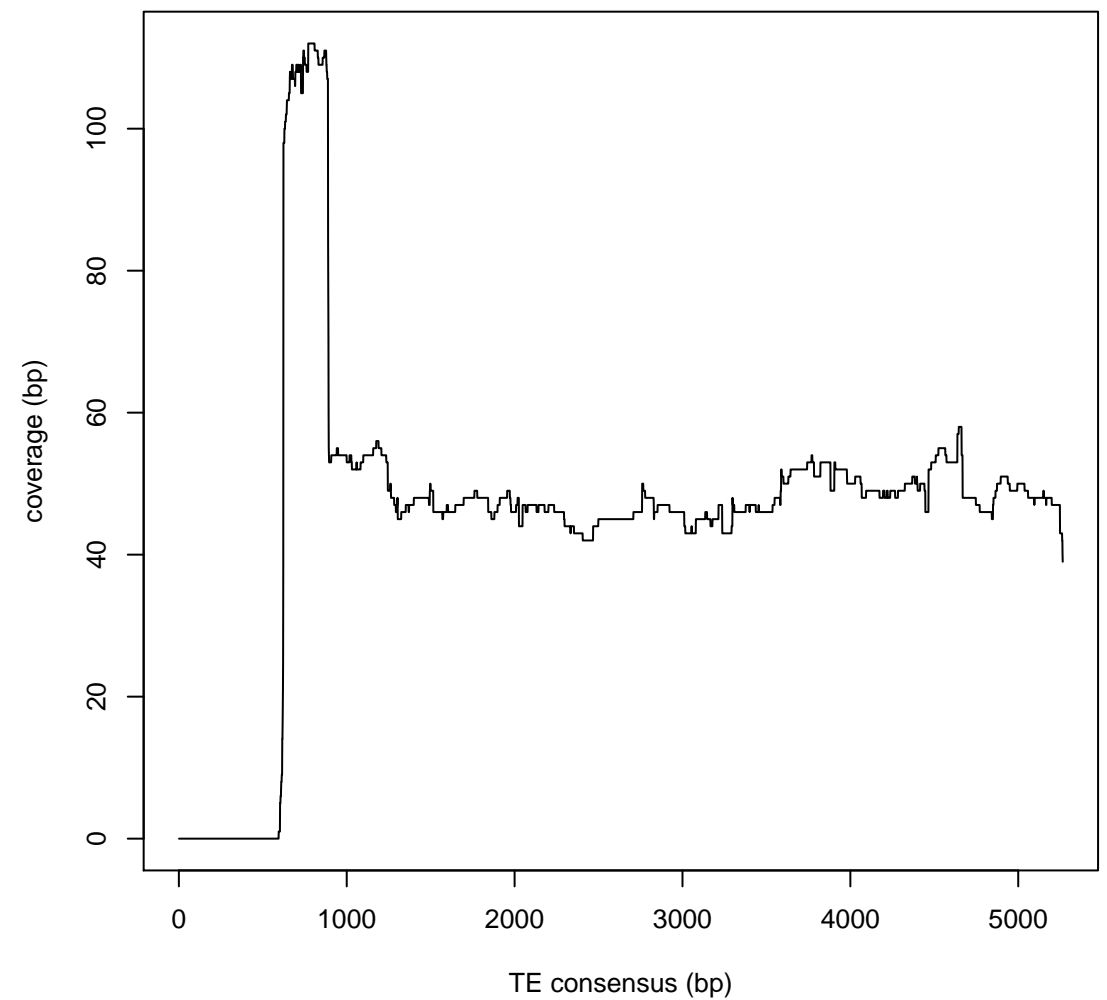


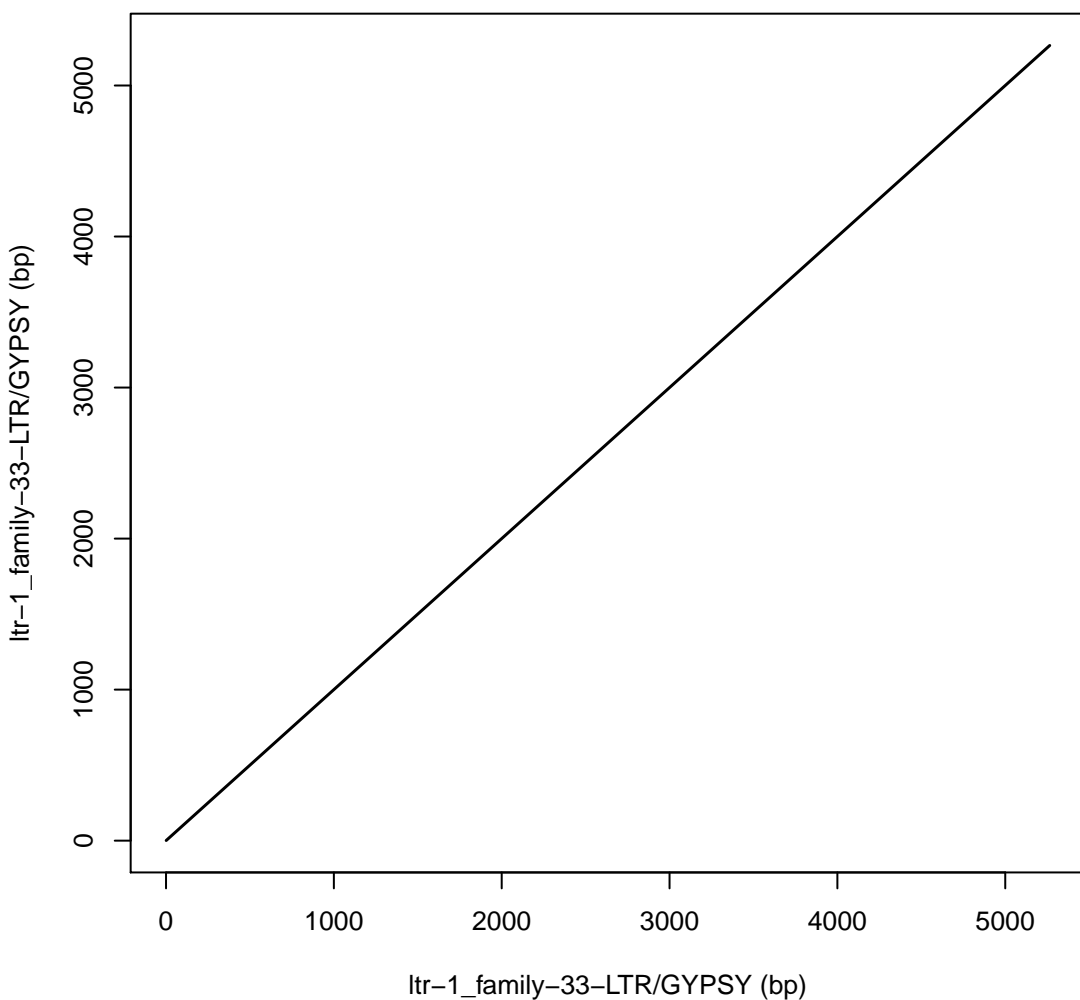
TE: ltr-1_family-33-LTR/GYPSY
consensus size: 5265bp; fragments: 243; full length: 0 (>=4738.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

