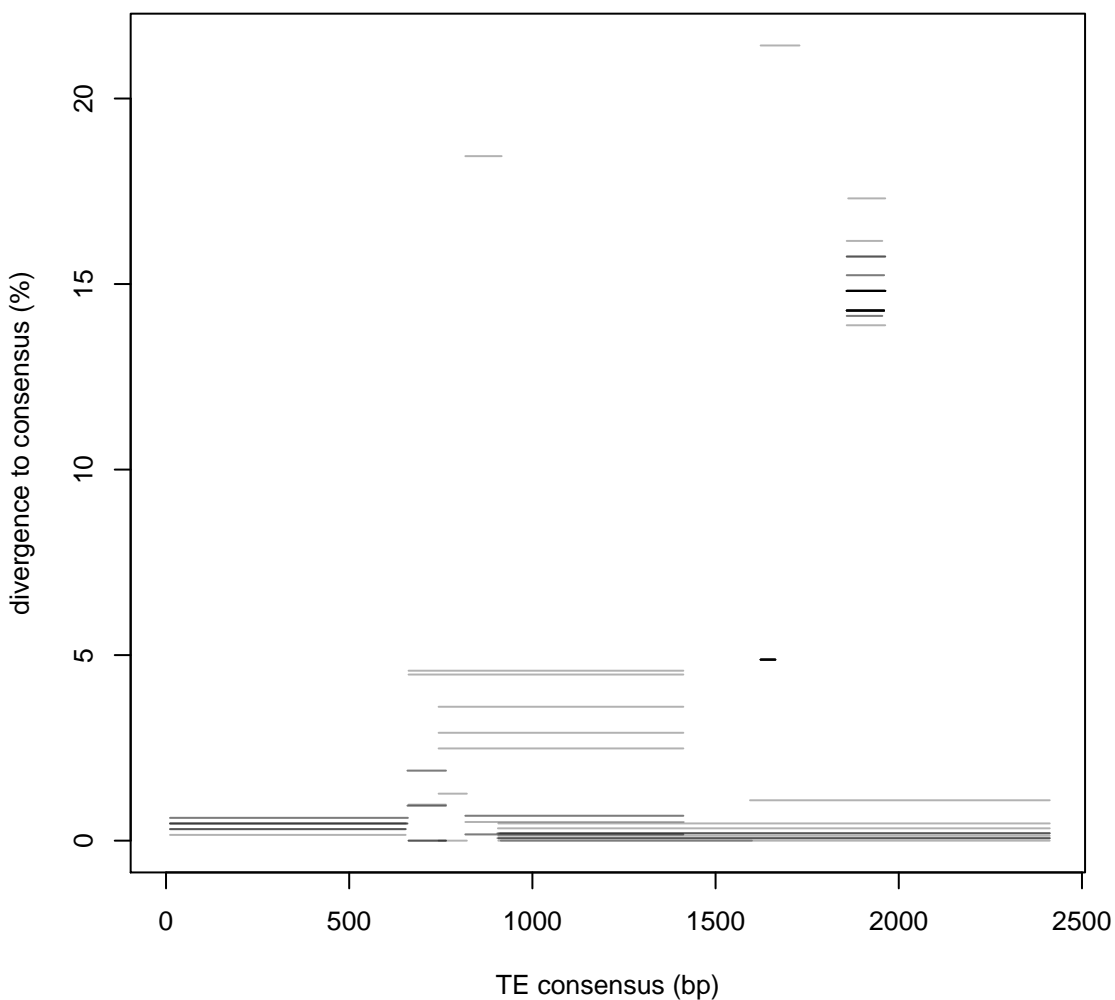
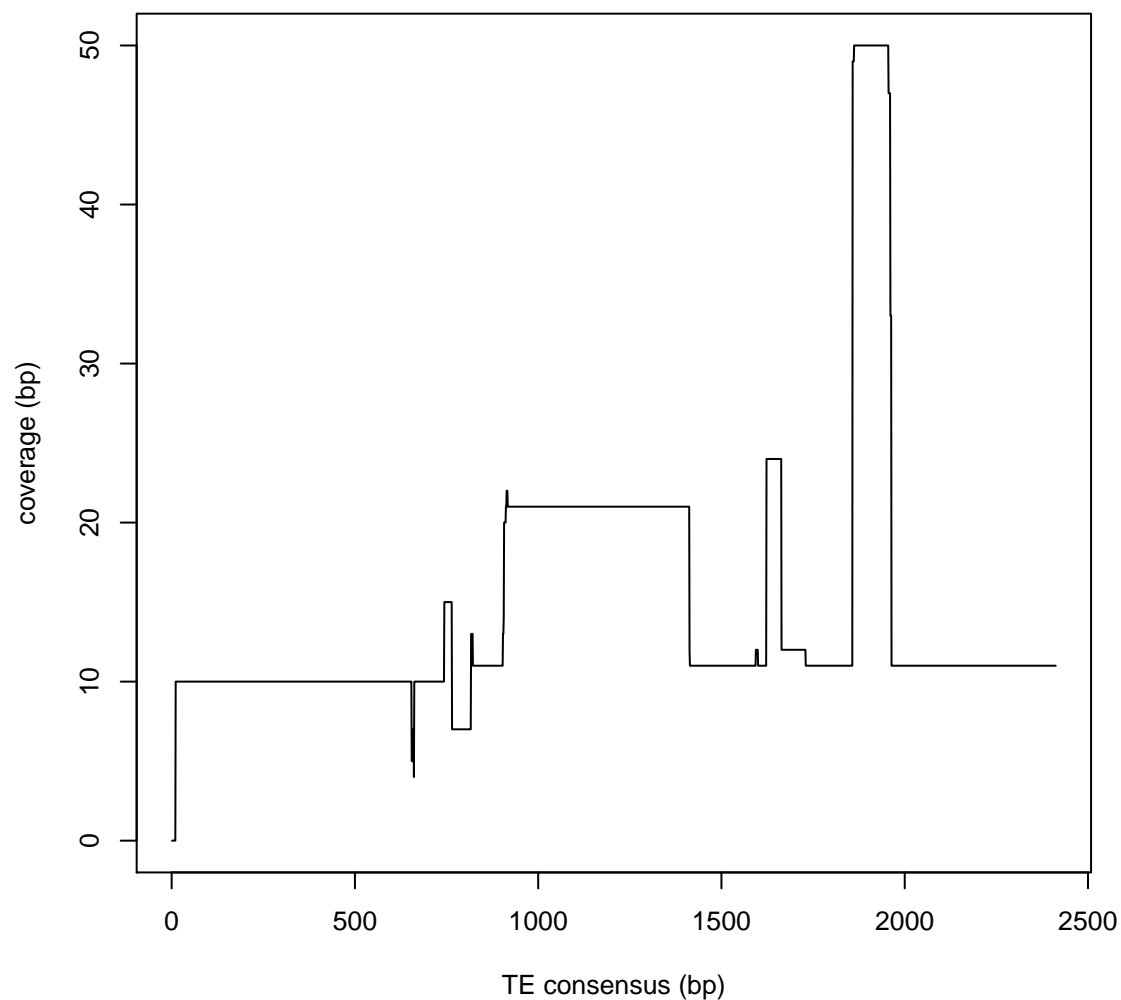


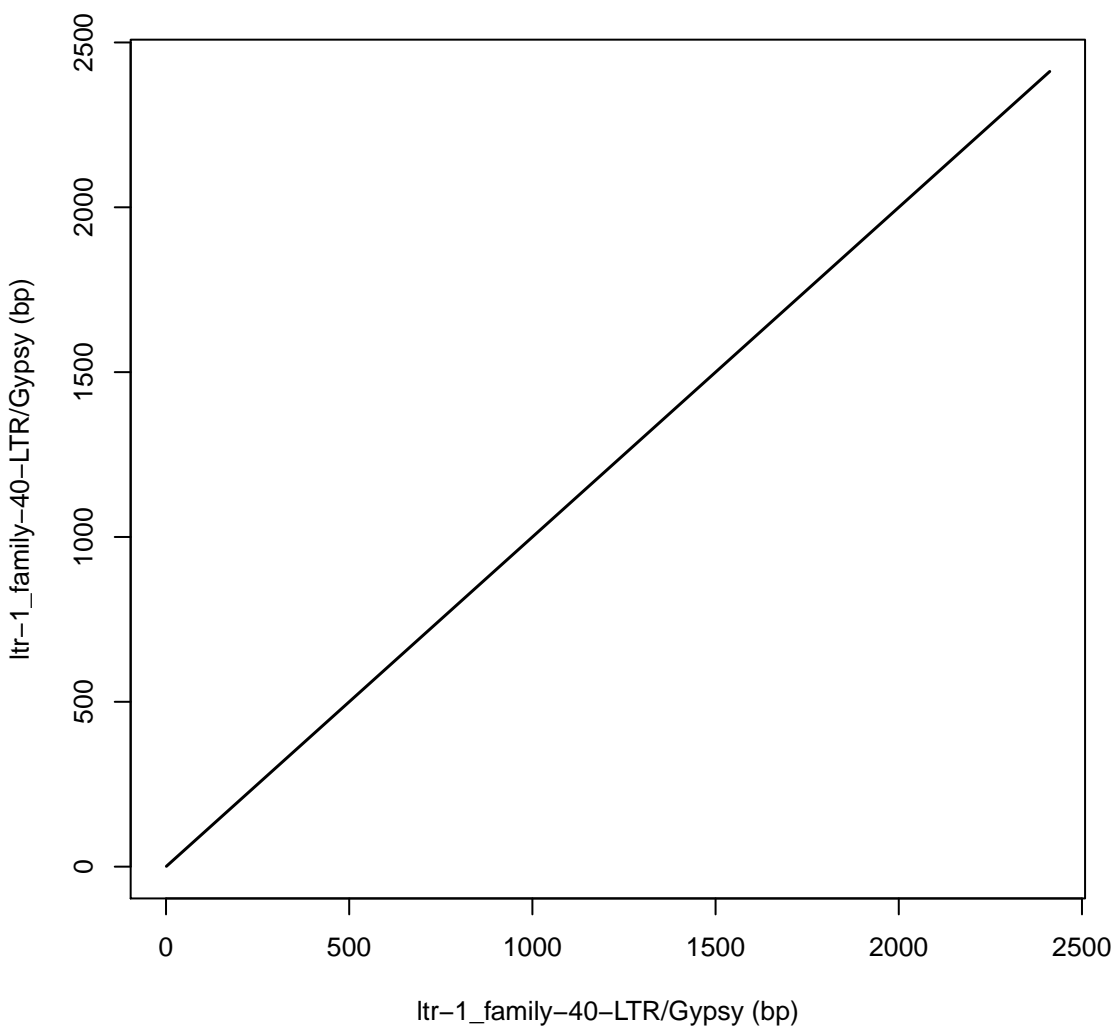
TE: ltr-1_family-40-LTR/Gypsy
consensus size: 2412bp; fragments: 95; full length: 0 (≥ 2170.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

