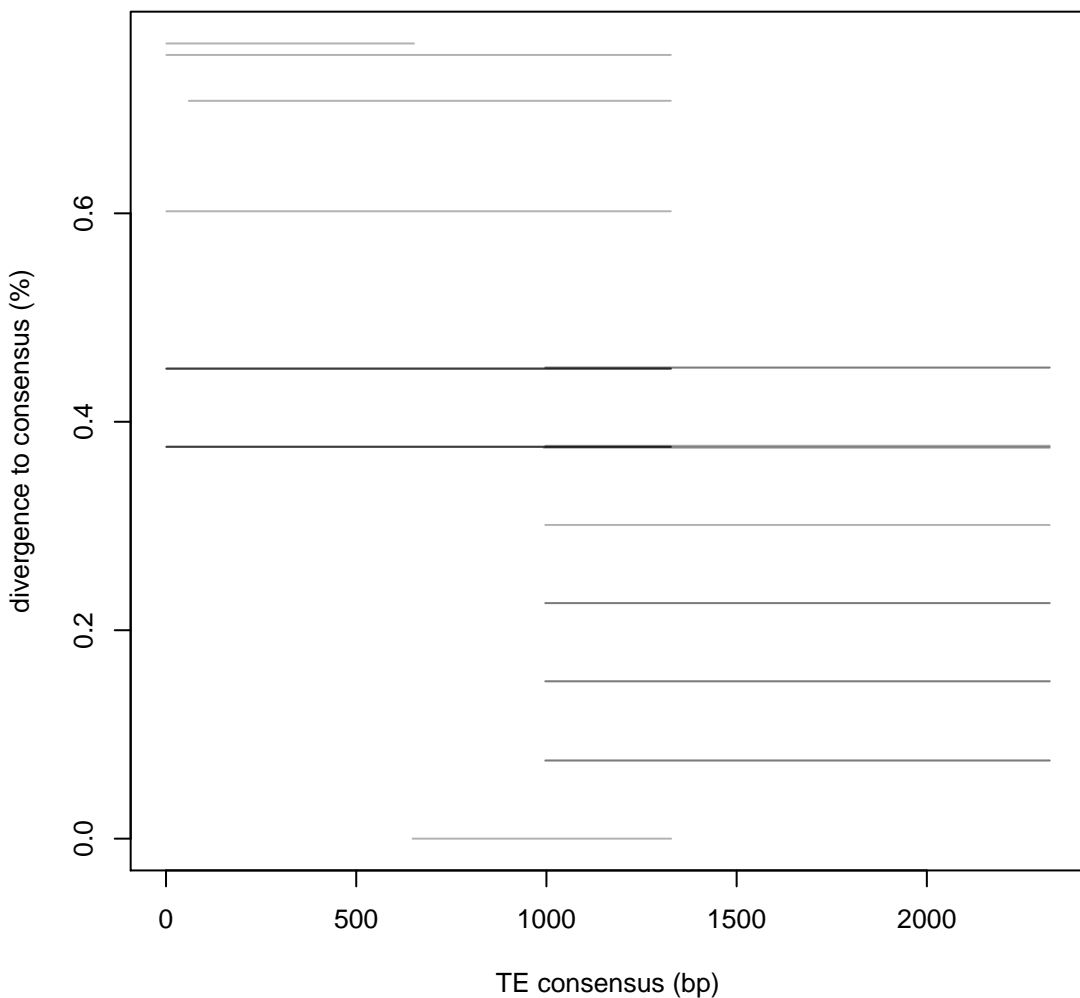
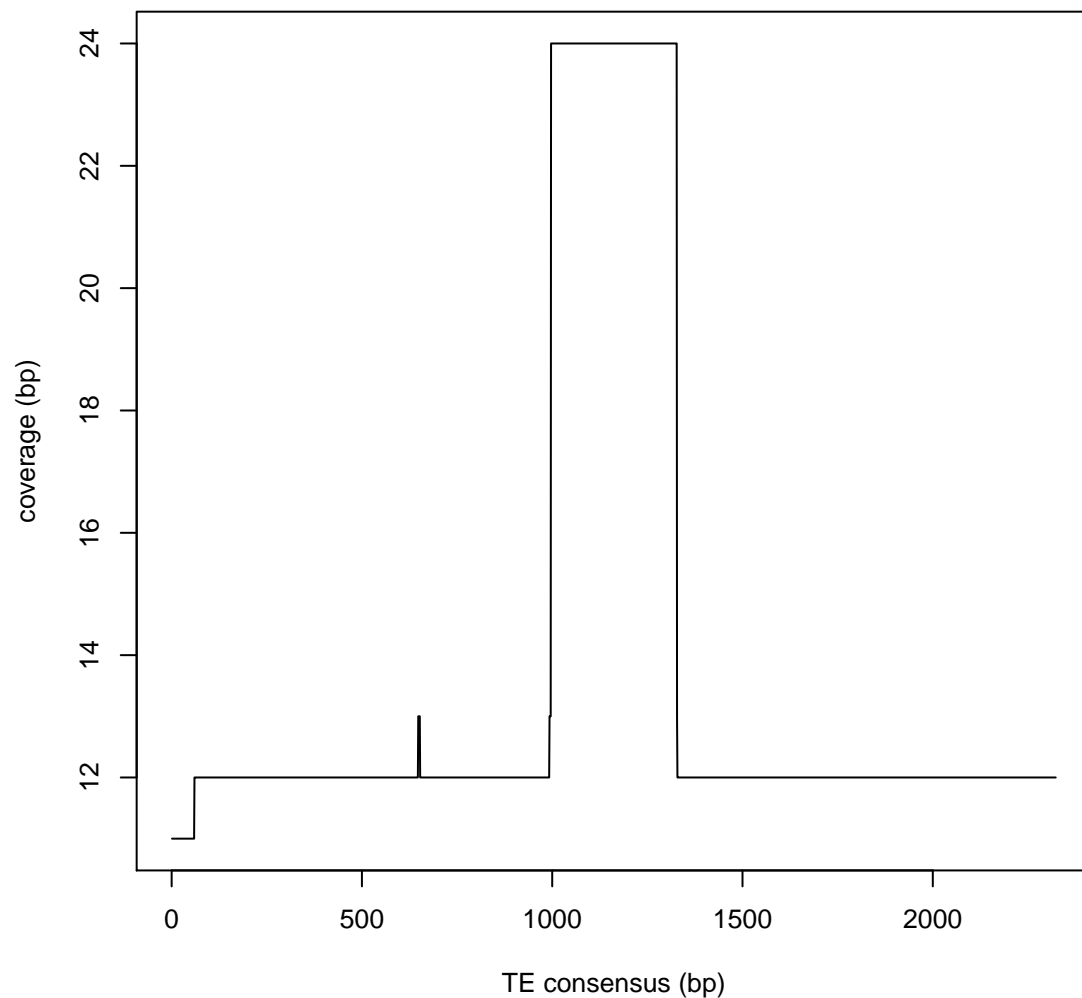


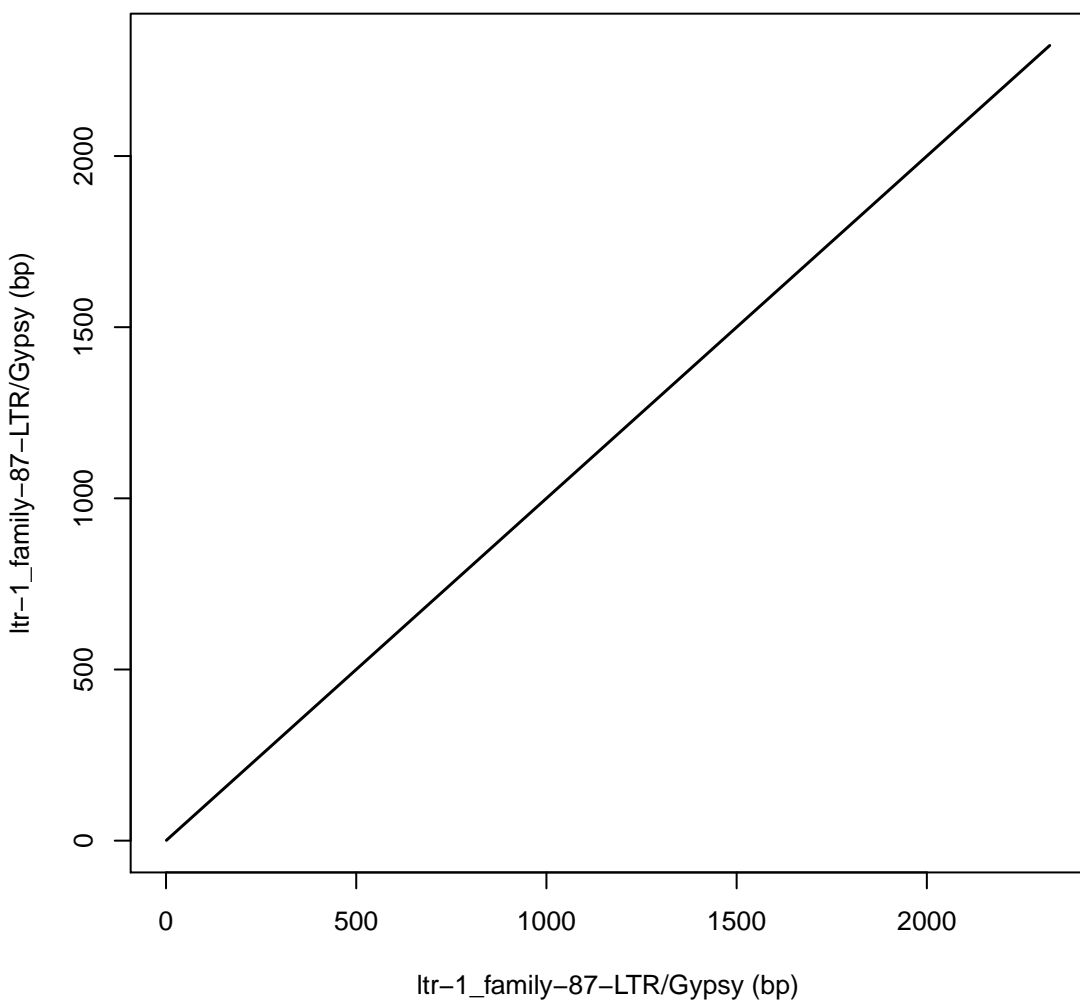
TE: ltr-1_family-87-LTR/Gypsy
consensus size: 2323bp; fragments: 25; full length: 0 (≥ 2090.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

