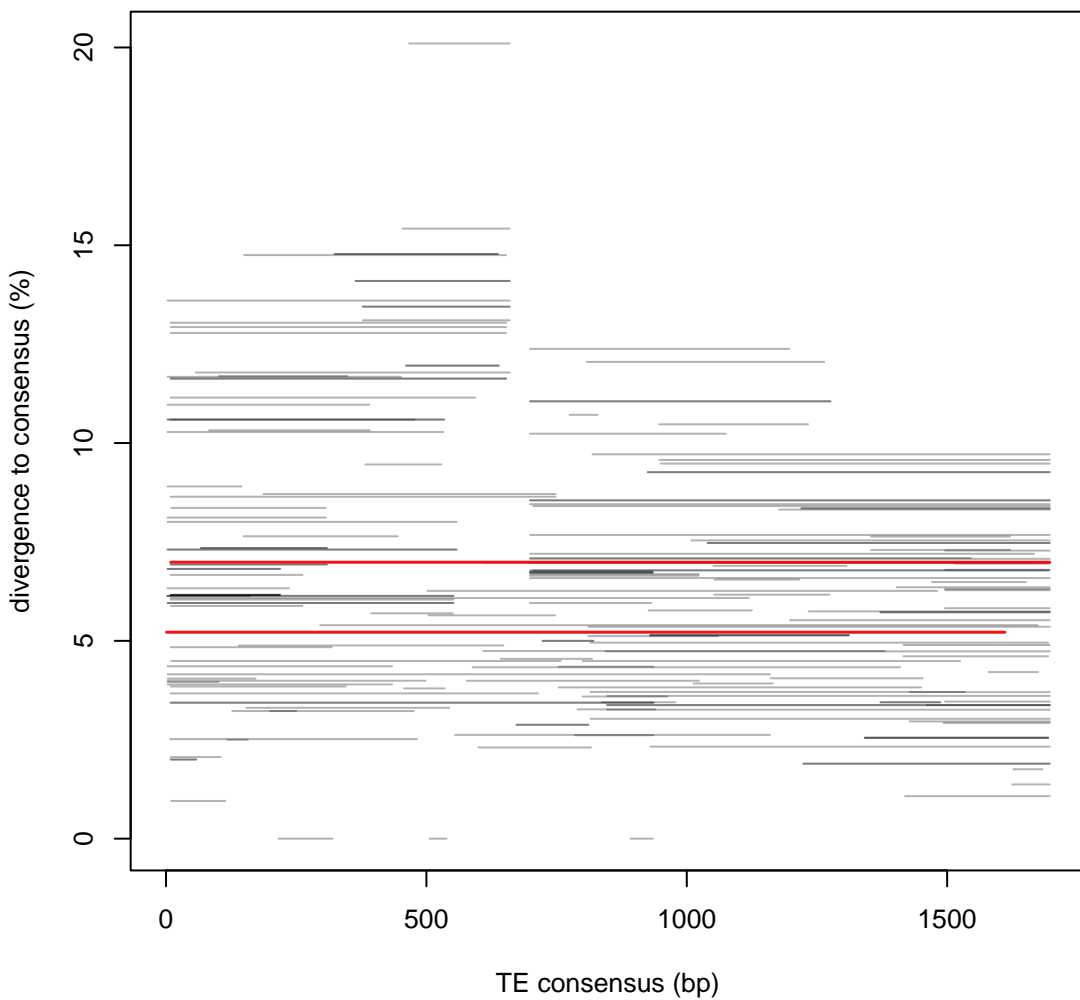
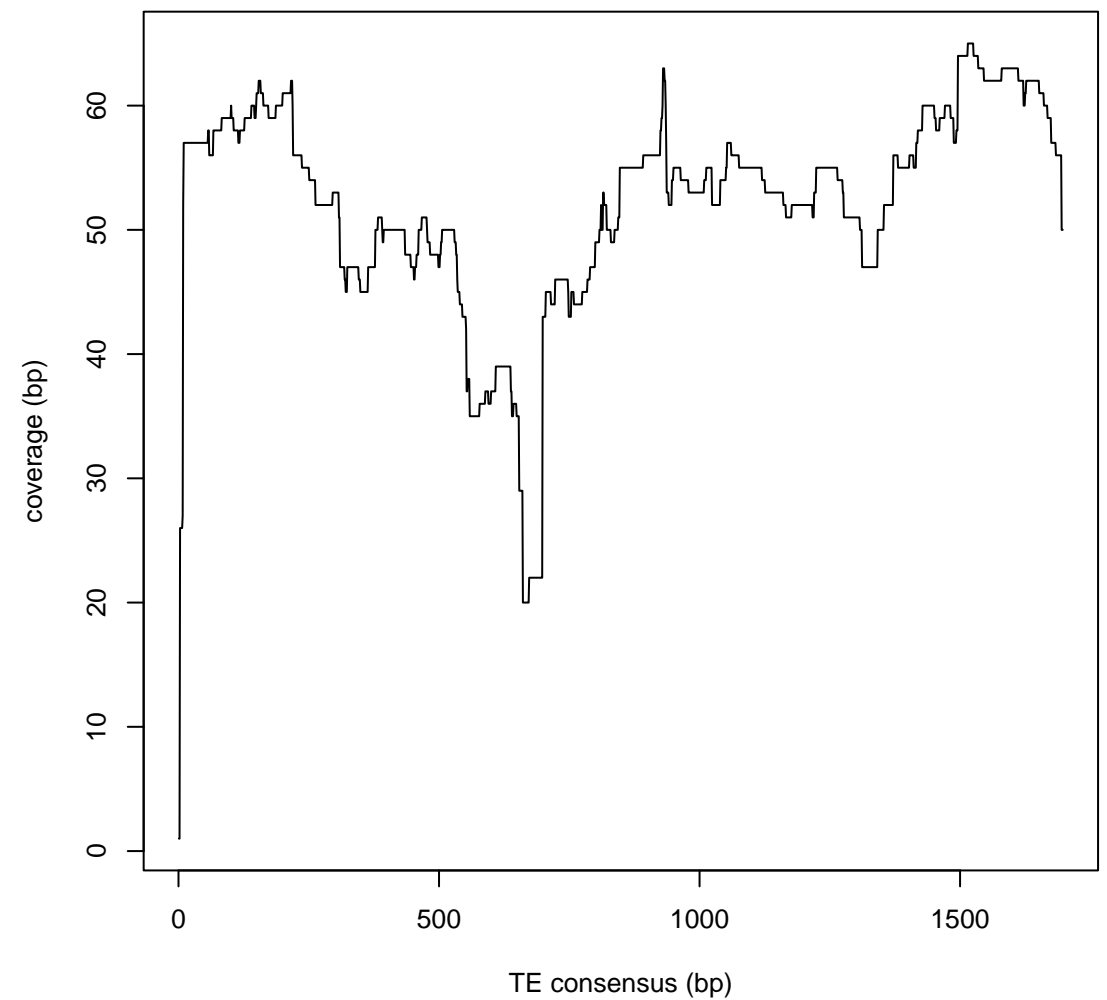


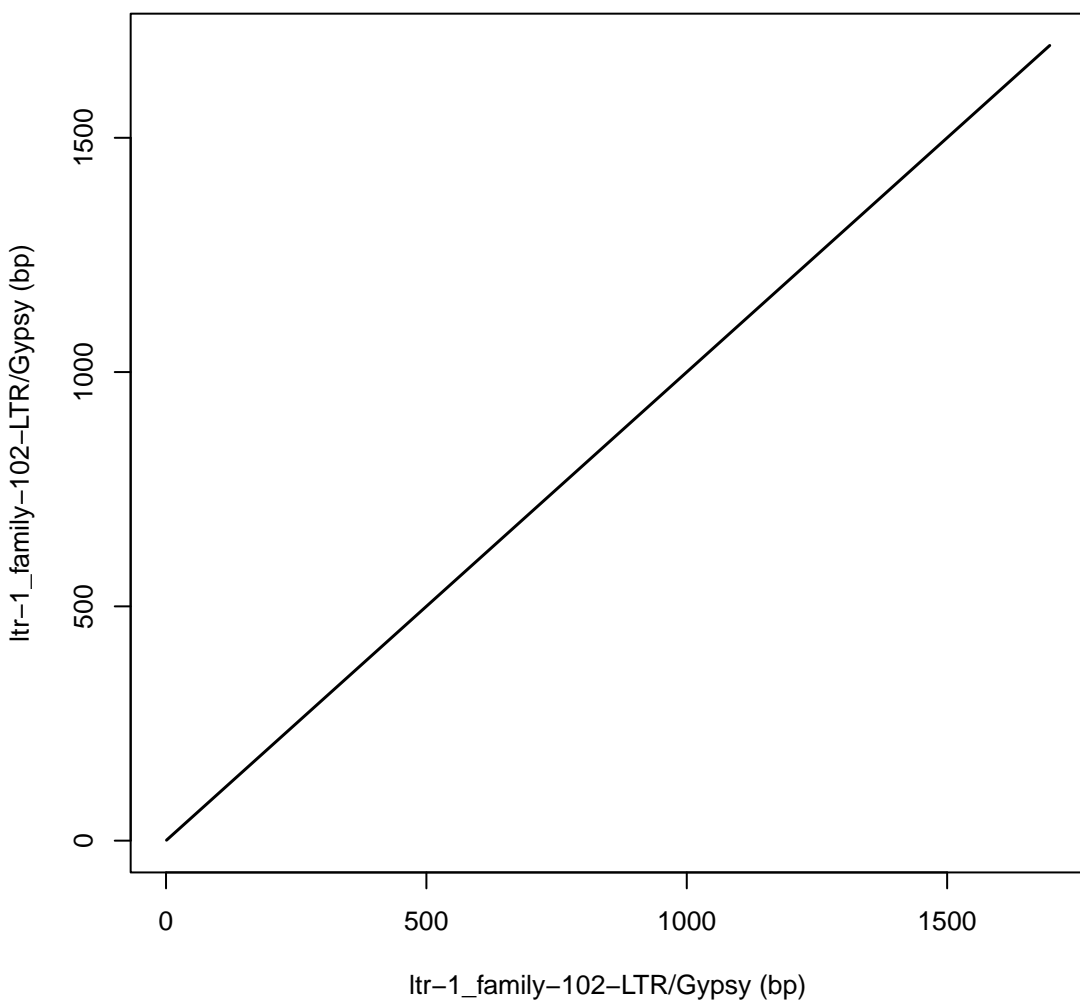
TE: ltr-1_family-102-LTR/Gypsy
consensus size: 1697bp; fragments: 197; full length: 2 (≥ 1527.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

