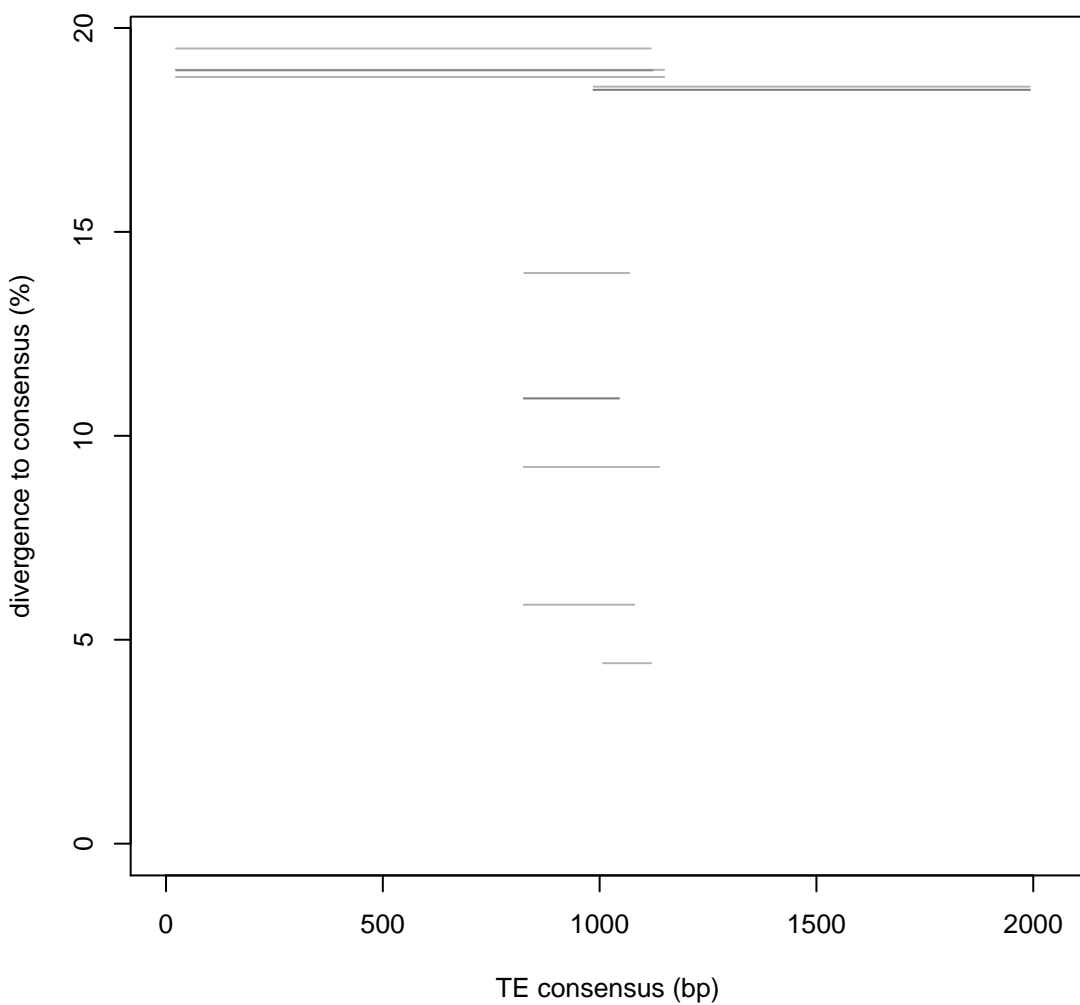
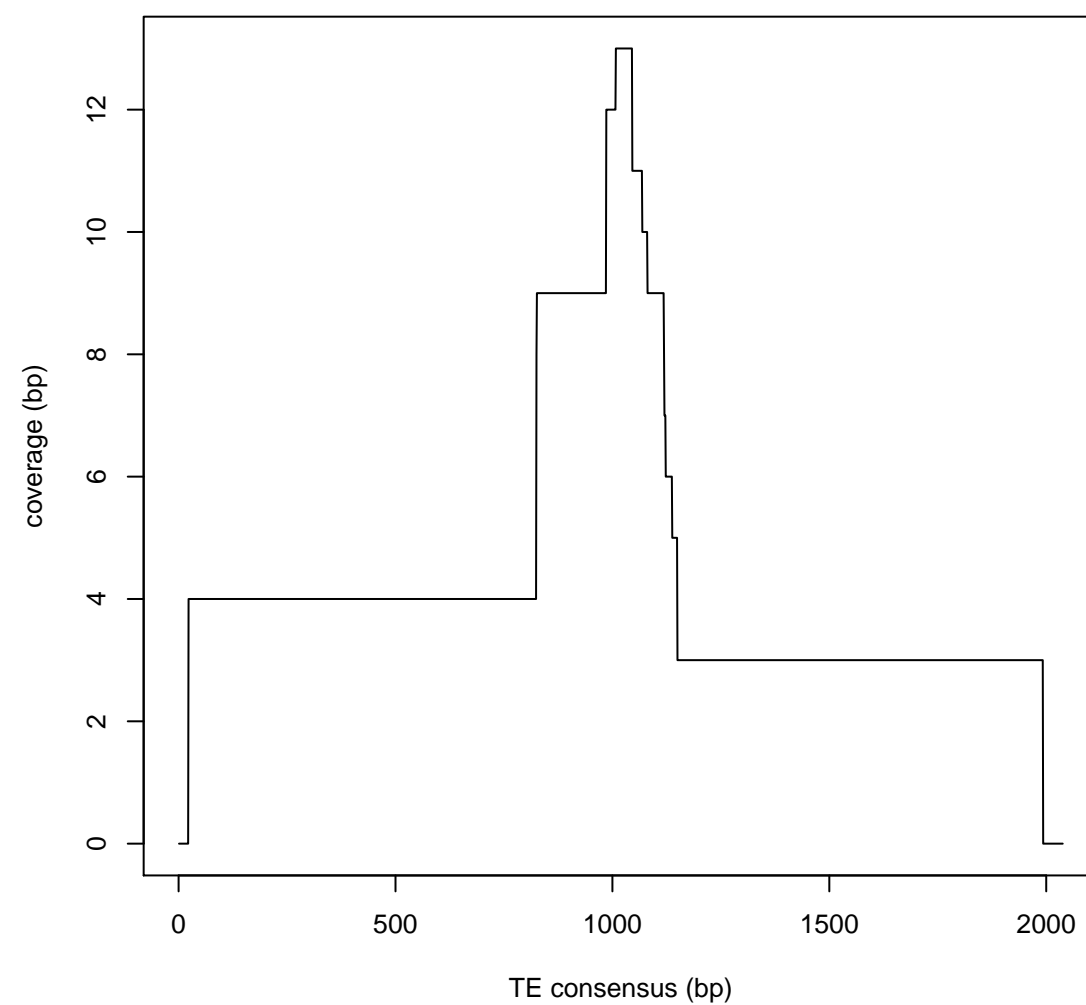


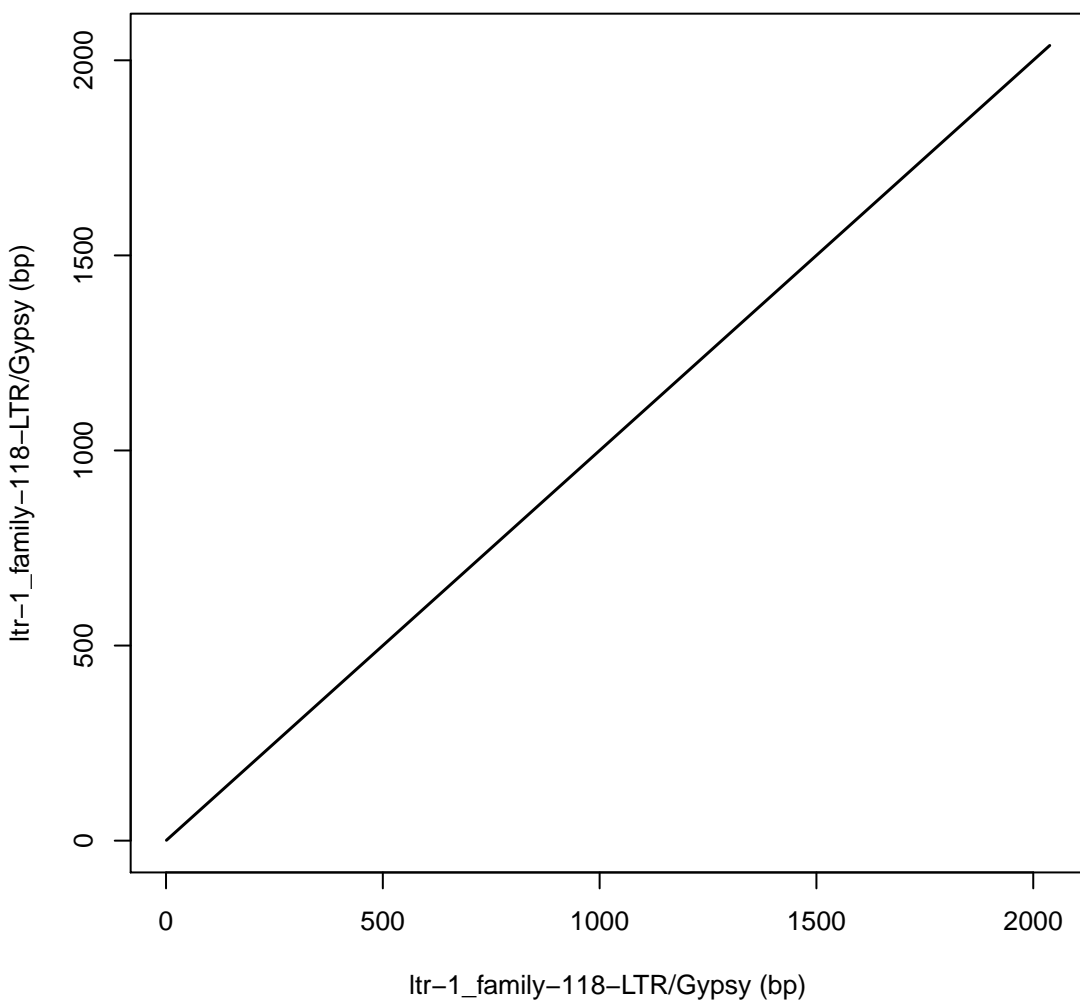
TE: ltr-1_family-118-LTR/Gypsy
consensus size: 2038bp; fragments: 13; full length: 0 (≥ 1834.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

