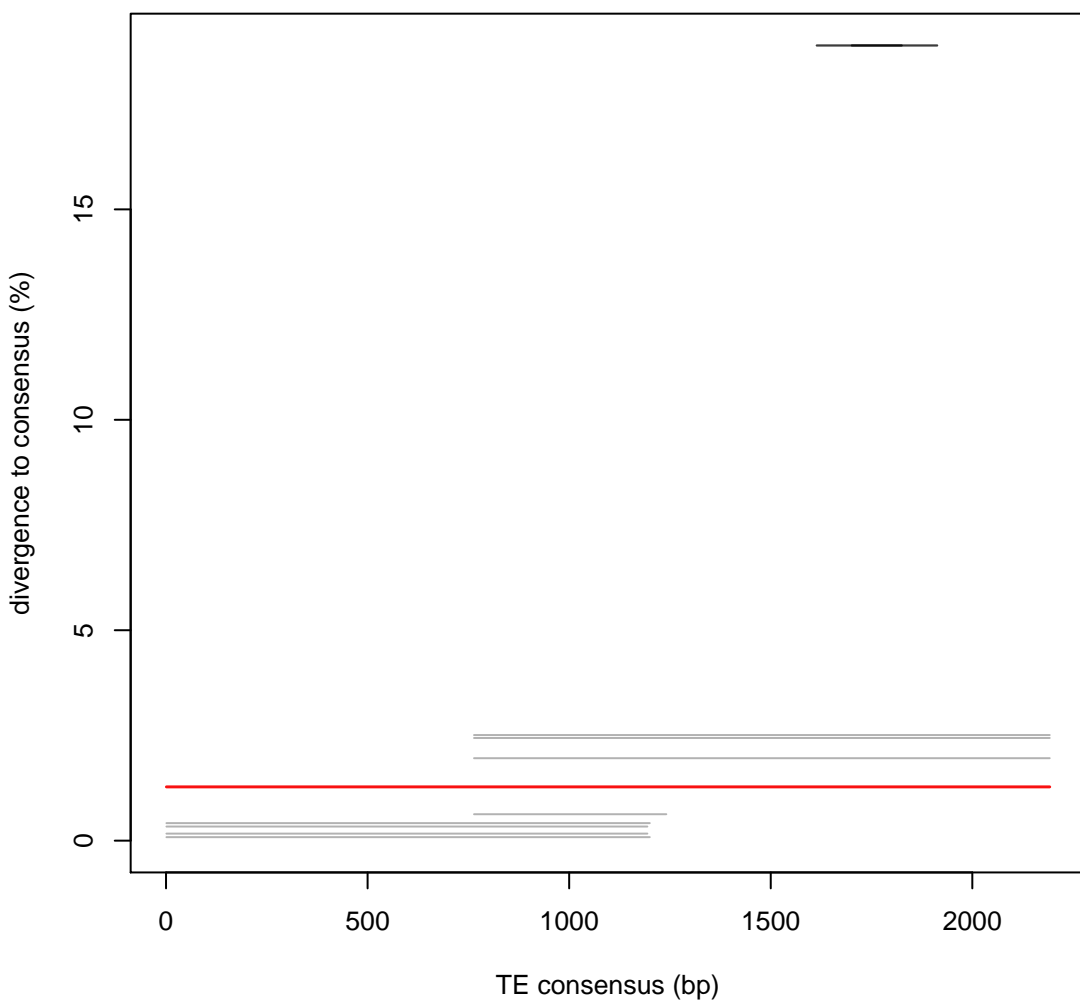
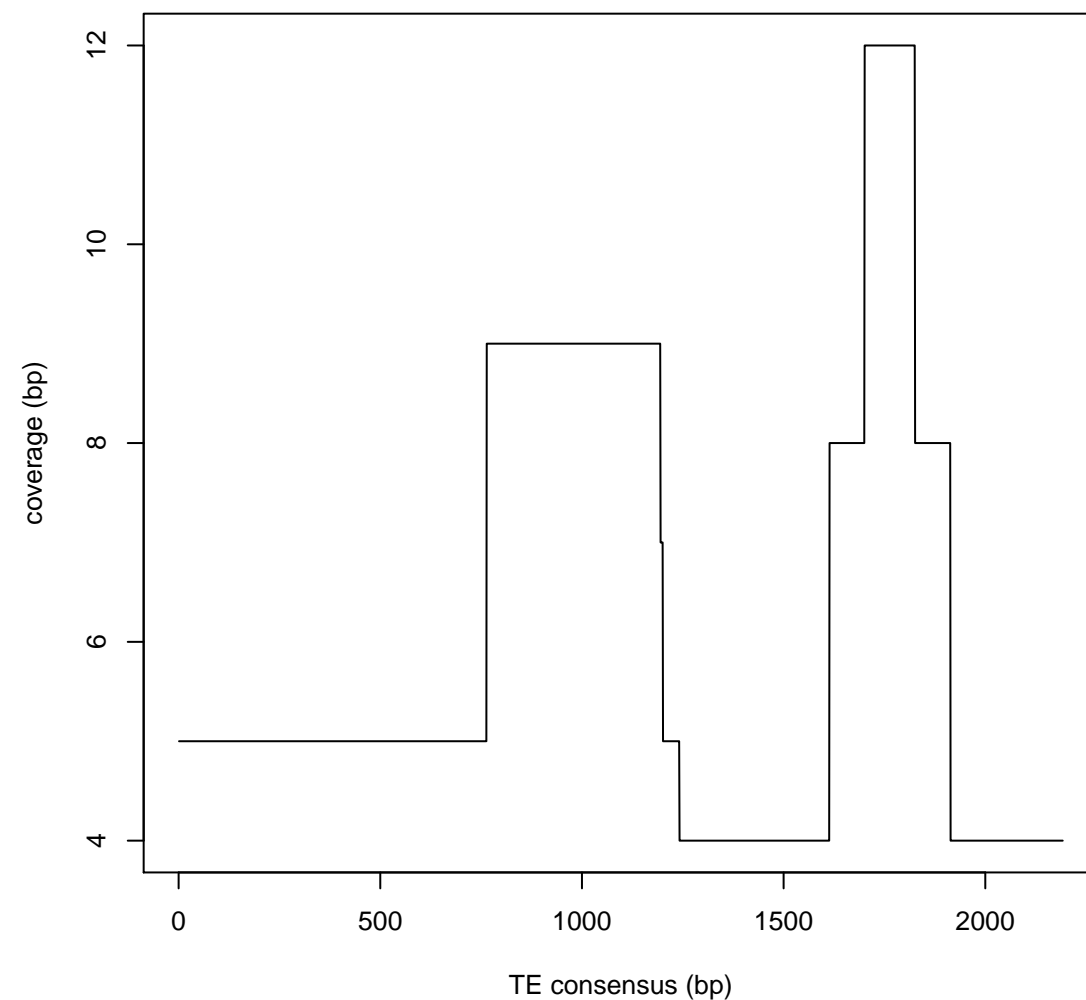


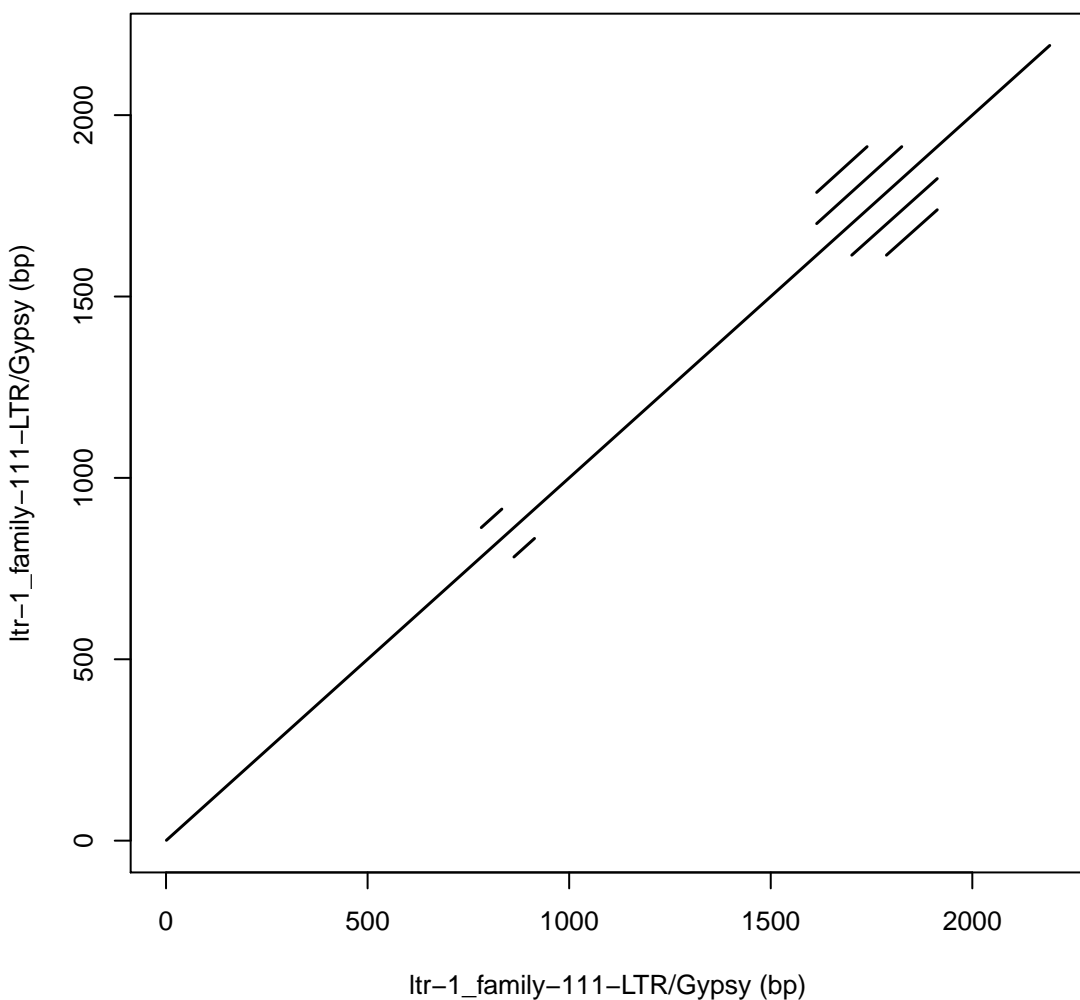
TE: ltr-1_family-111-LTR/Gypsy
consensus size: 2192bp; fragments: 17; full length: 1 (≥ 1972.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

