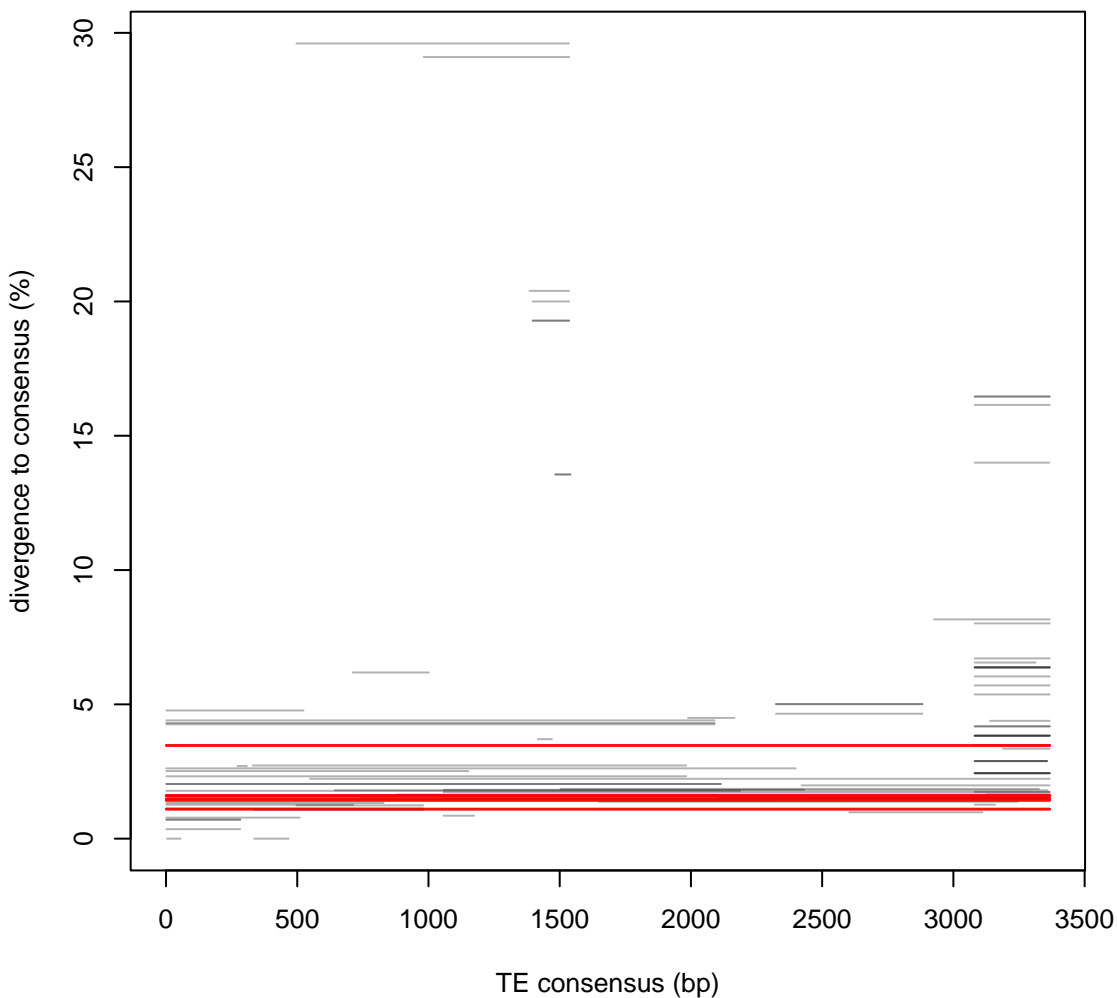
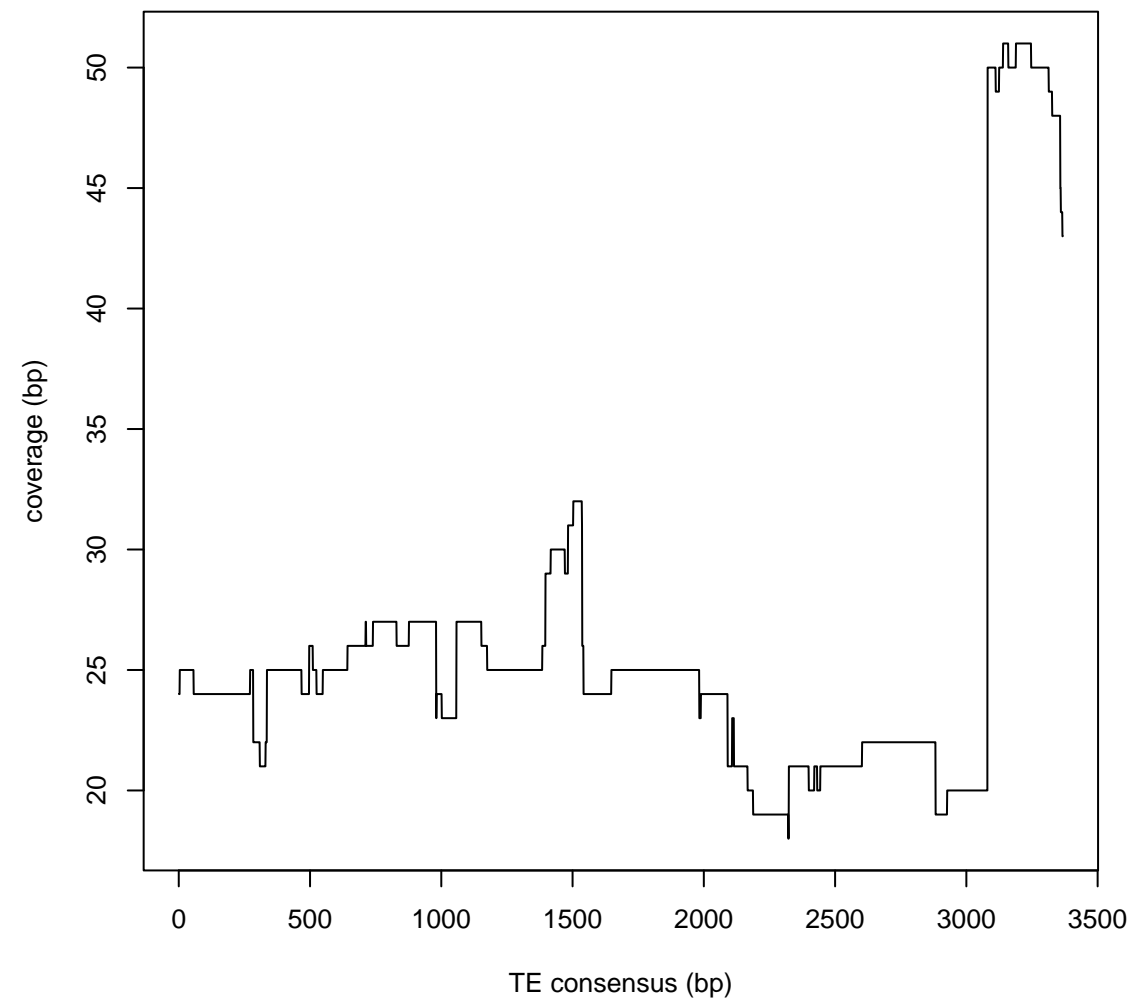


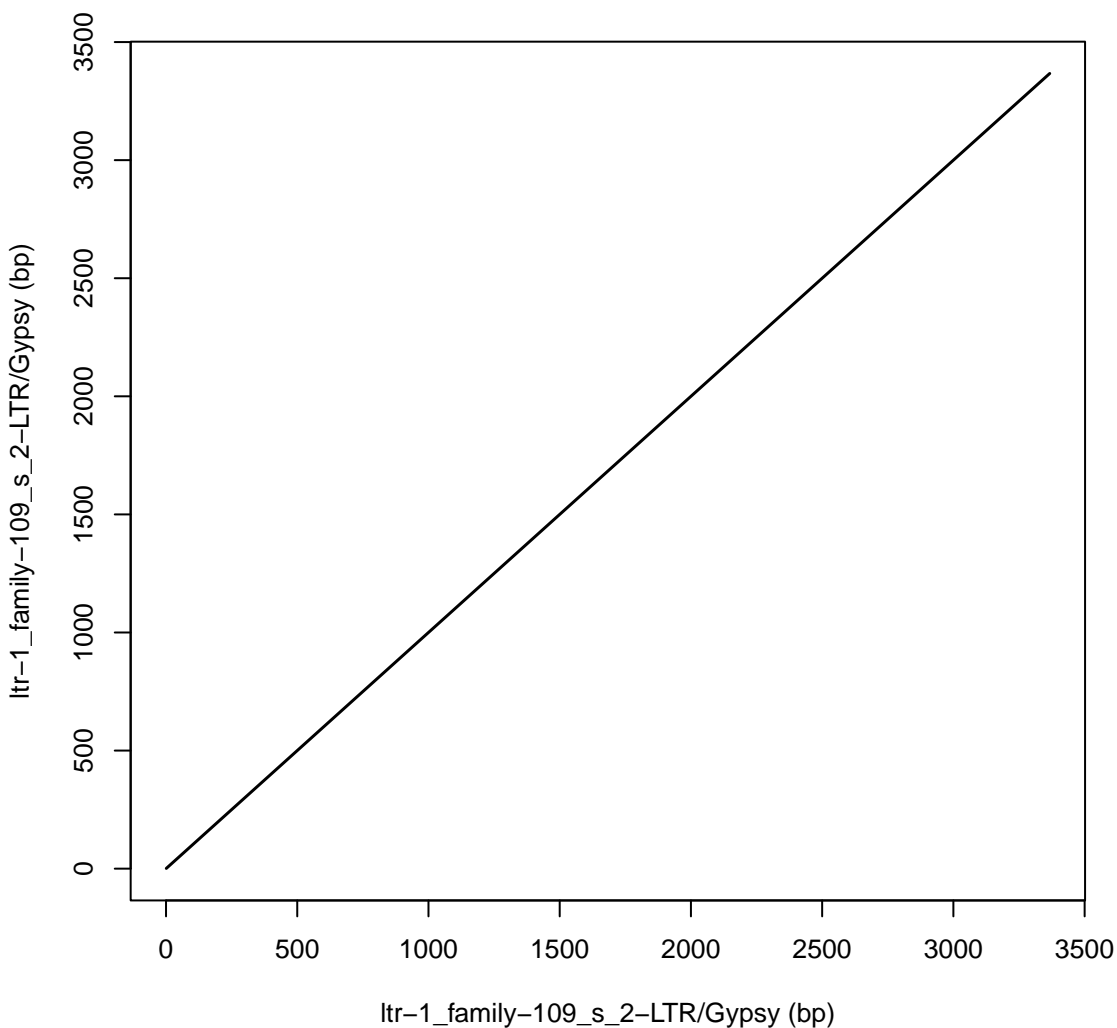
TE: ltr-1_family-109_s_2-LTR/Gypsy
consensus size: 3367bp; fragments: 94; full length: 6 (≥ 3030.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

