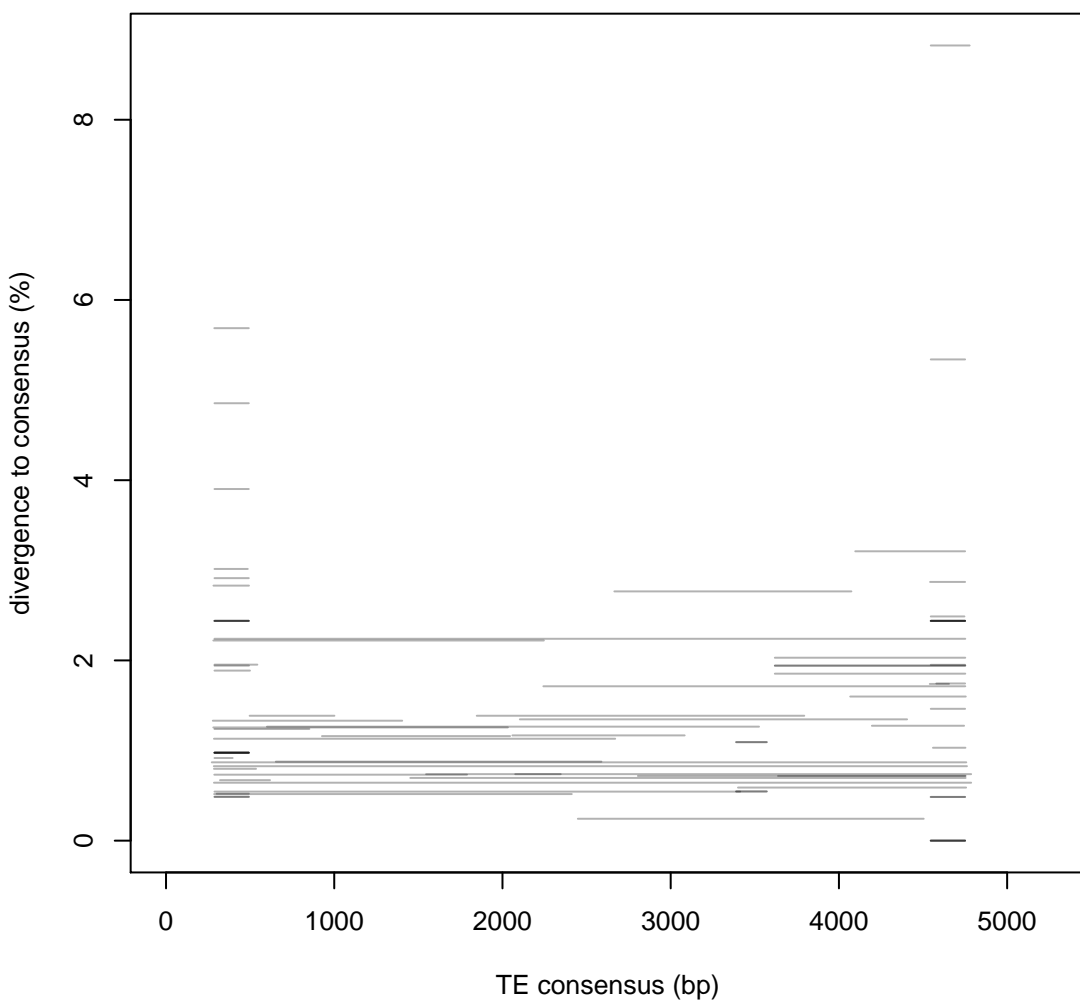
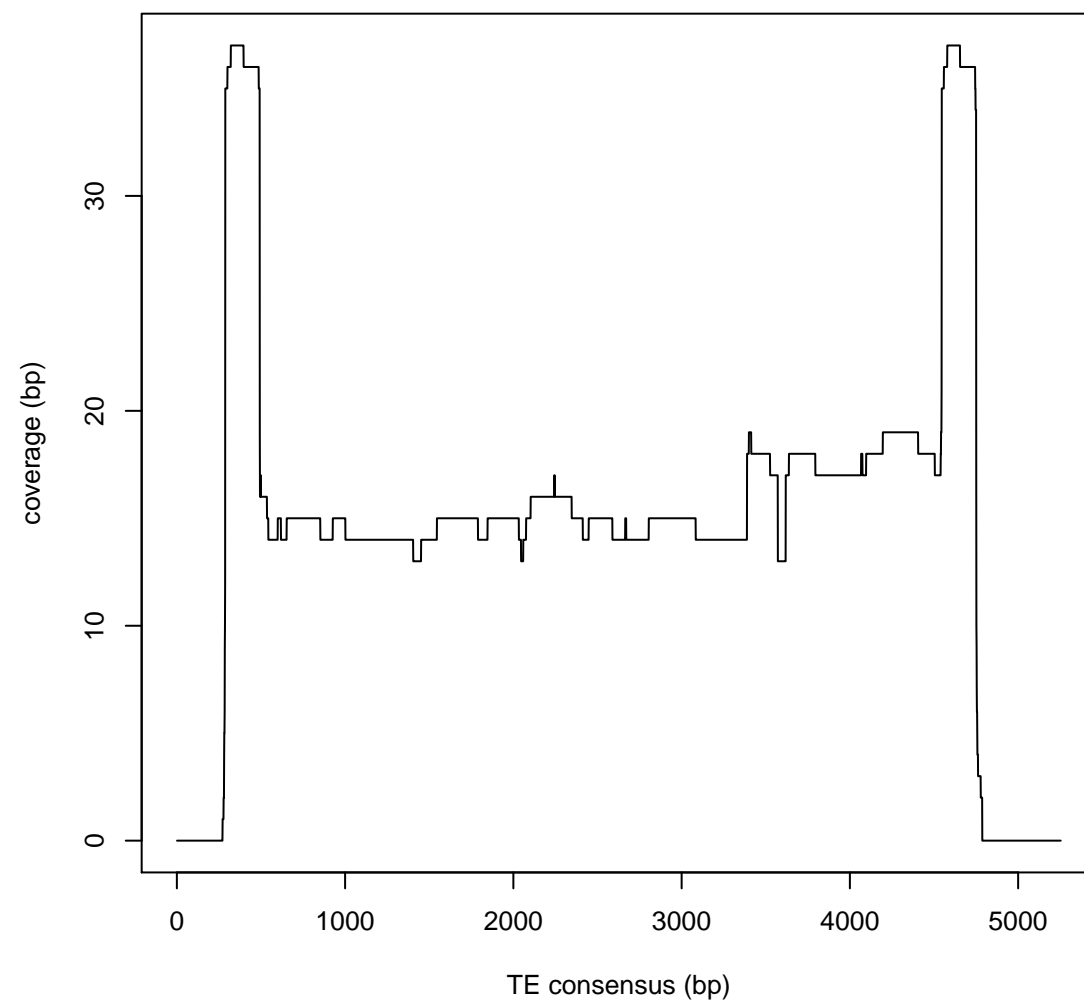


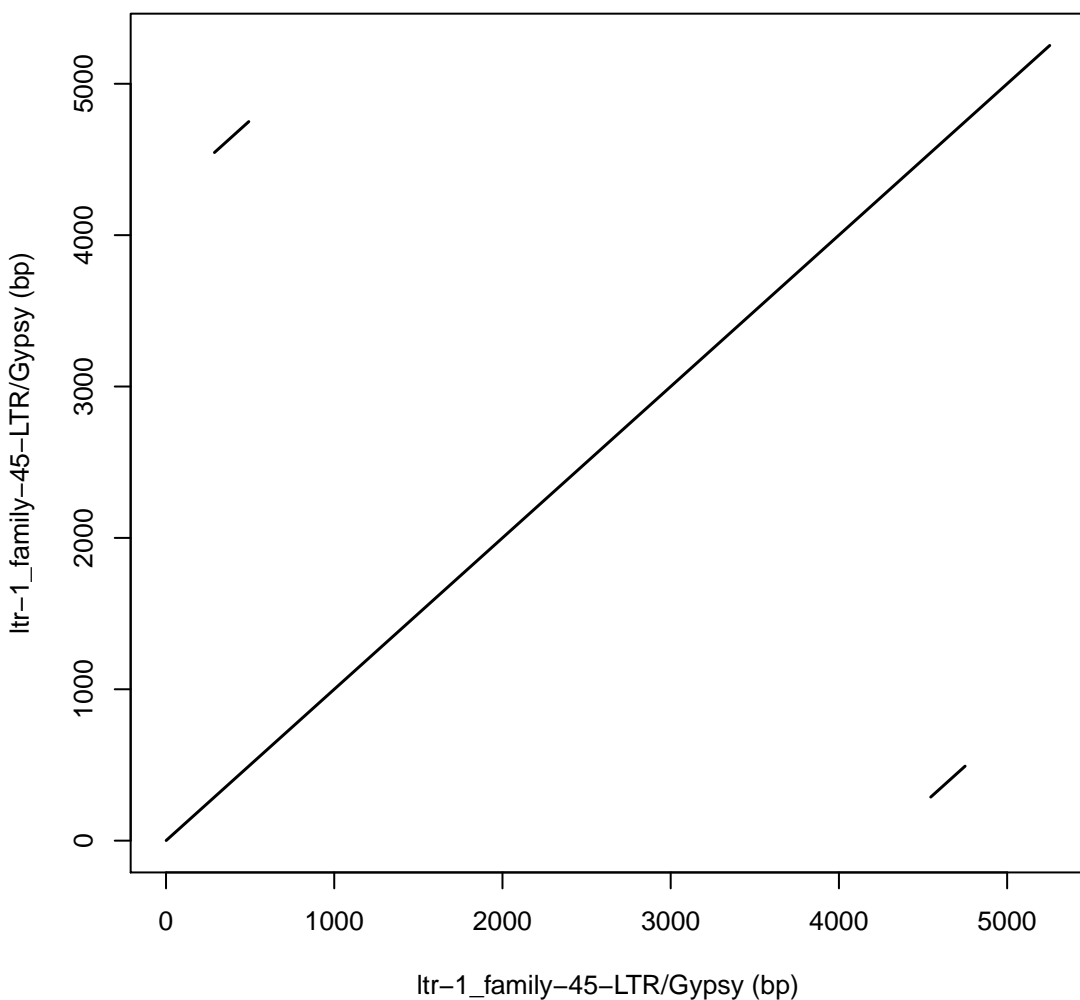
TE: ltr-1_family-45-LTR/Gypsy
consensus size: 5253bp; fragments: 84; full length: 0 (≥ 4727.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

