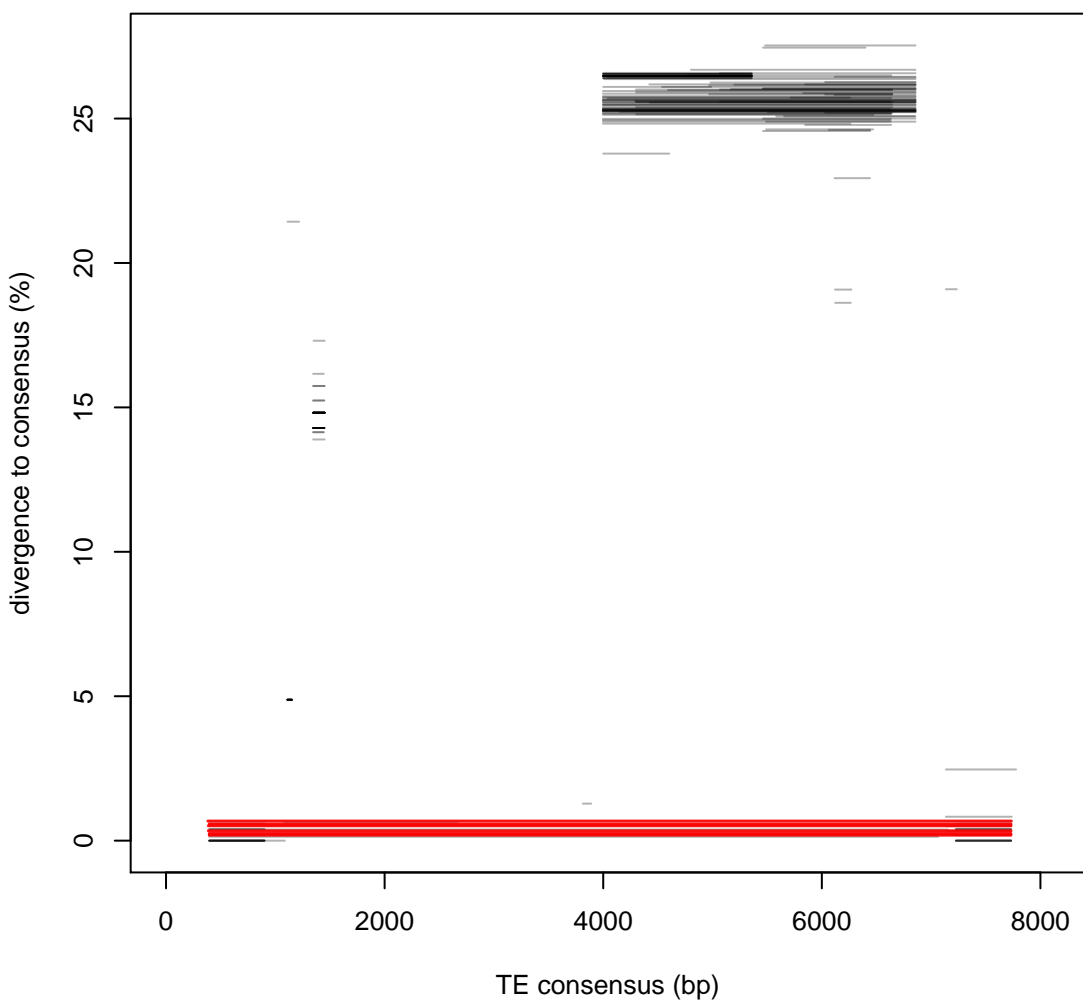
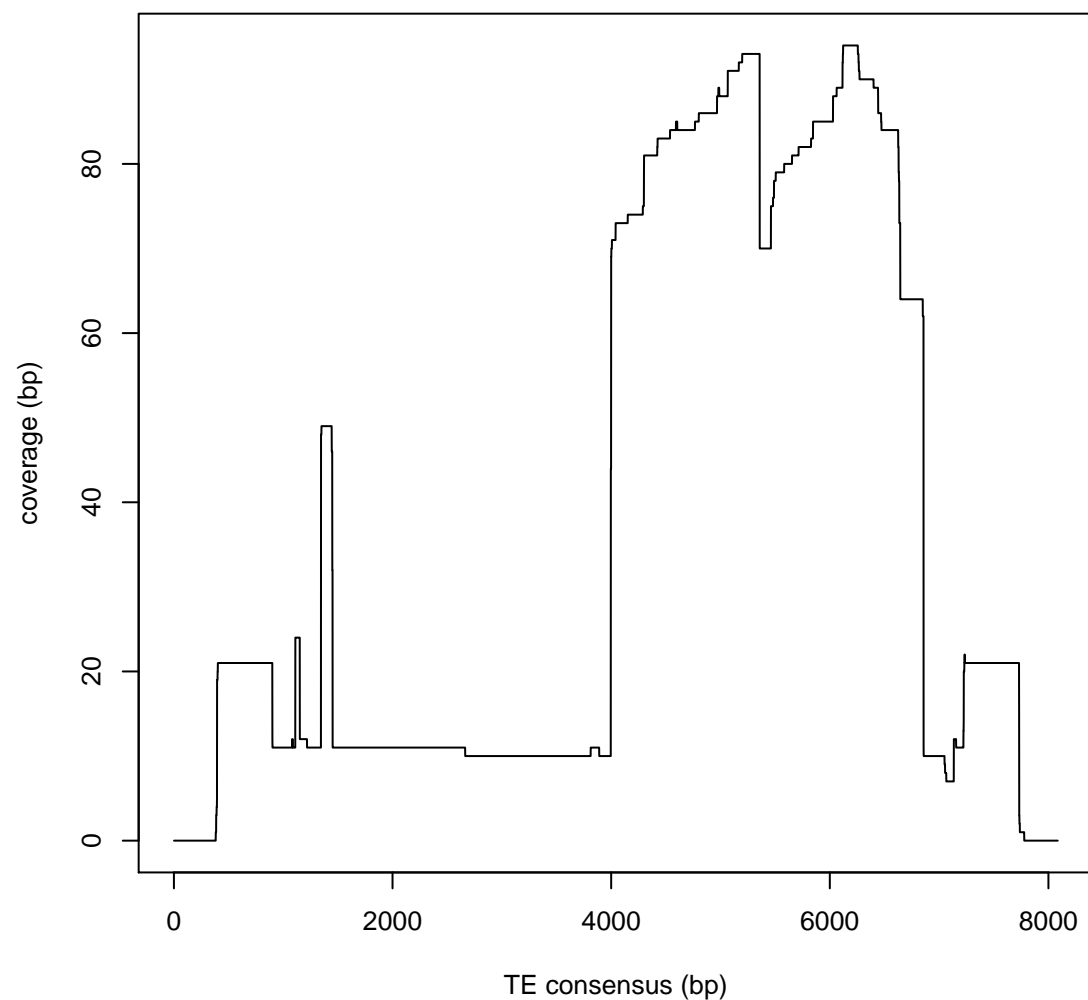


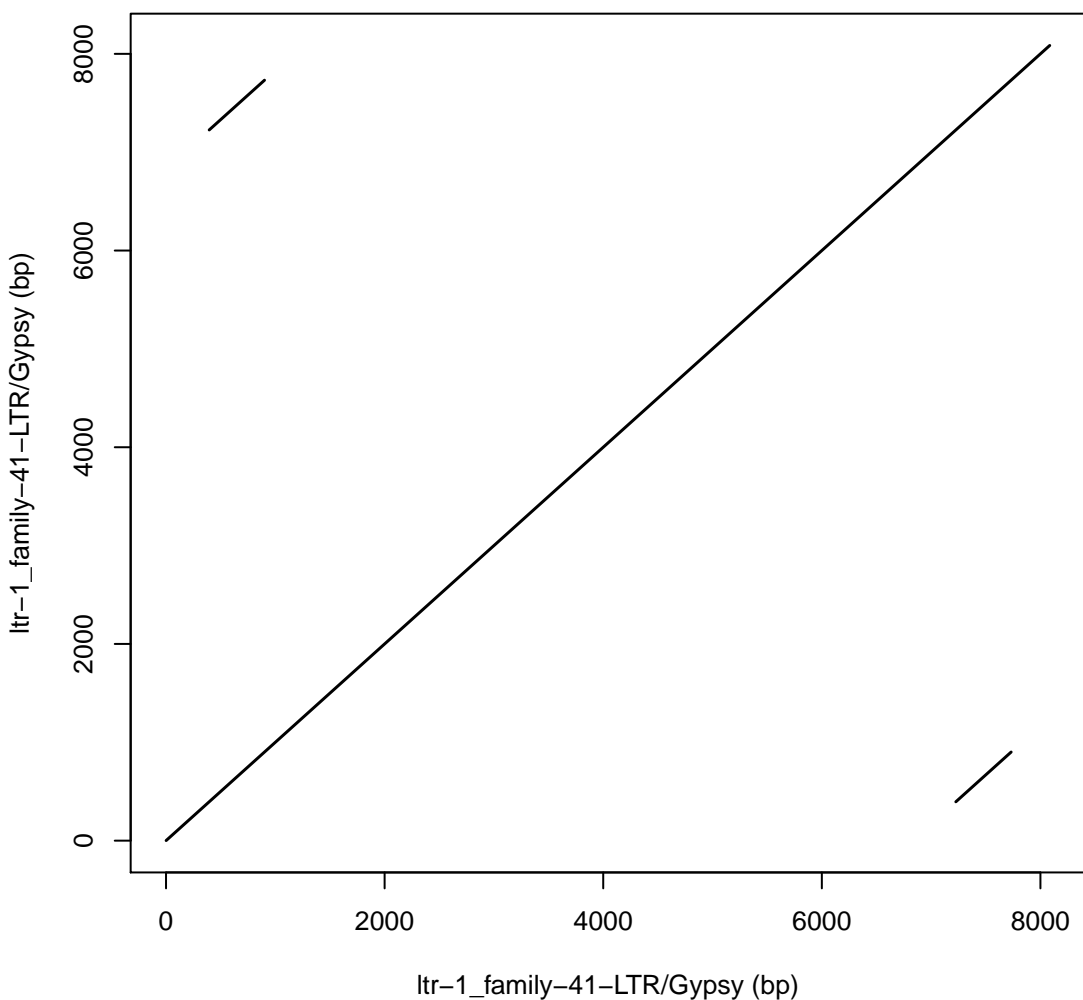
TE: ltr-1_family-41-LTR/Gypsy
consensus size: 8085bp; fragments: 199; full length: 6 (≥ 7276.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

