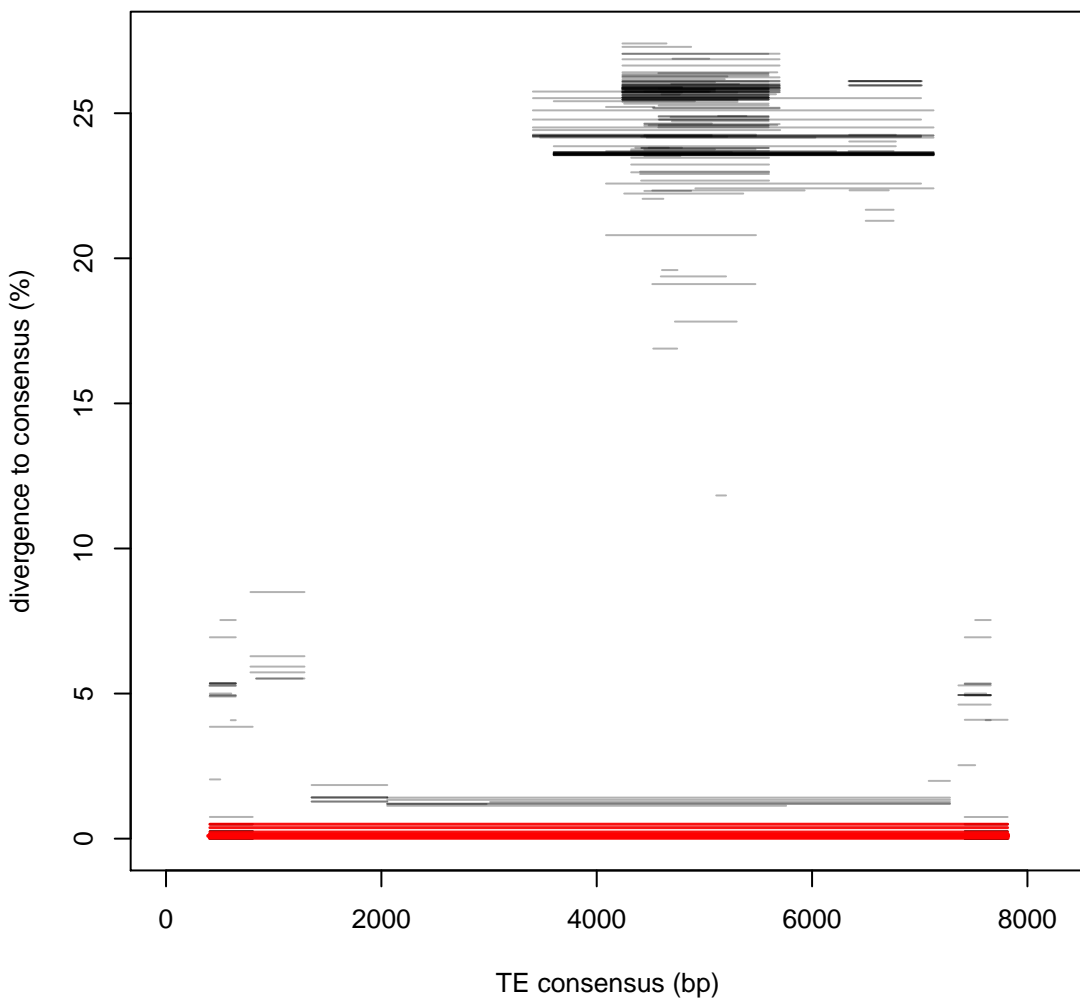
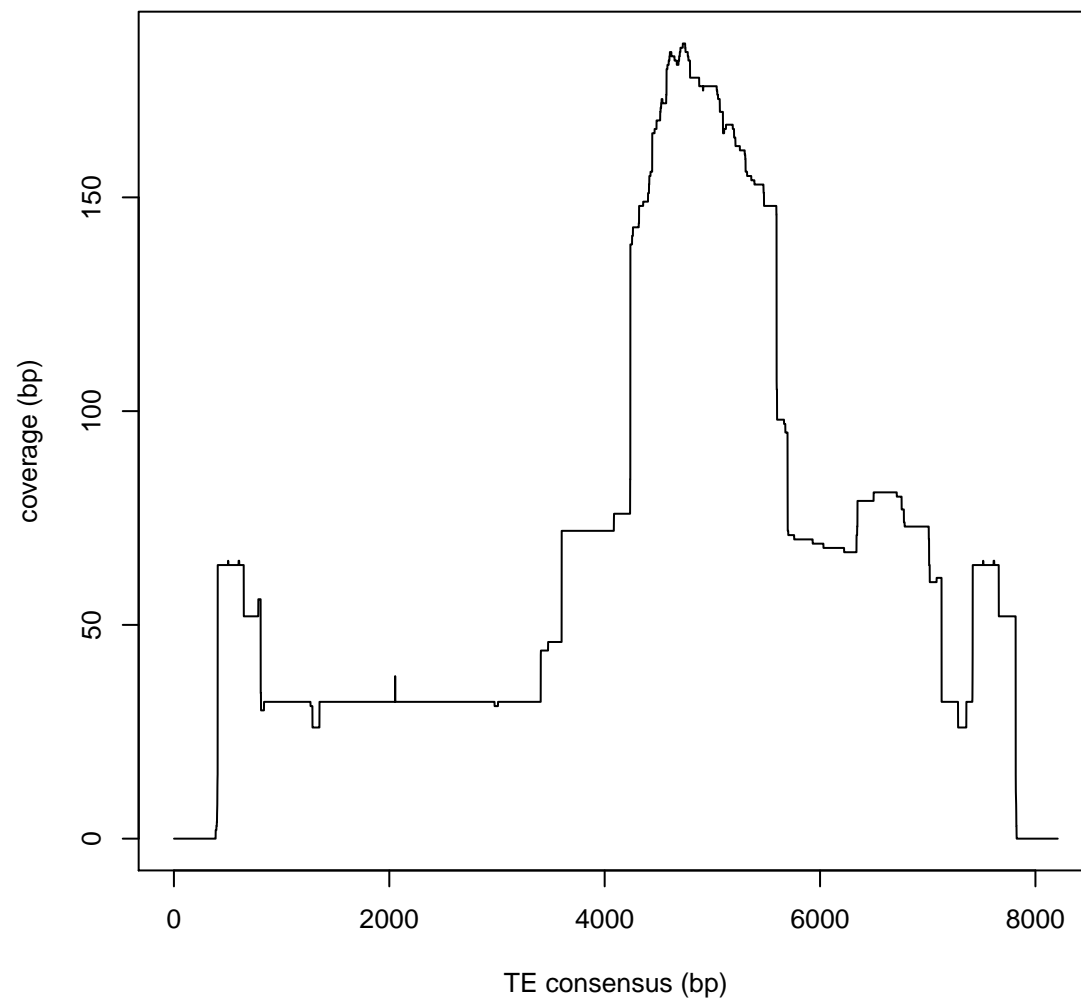


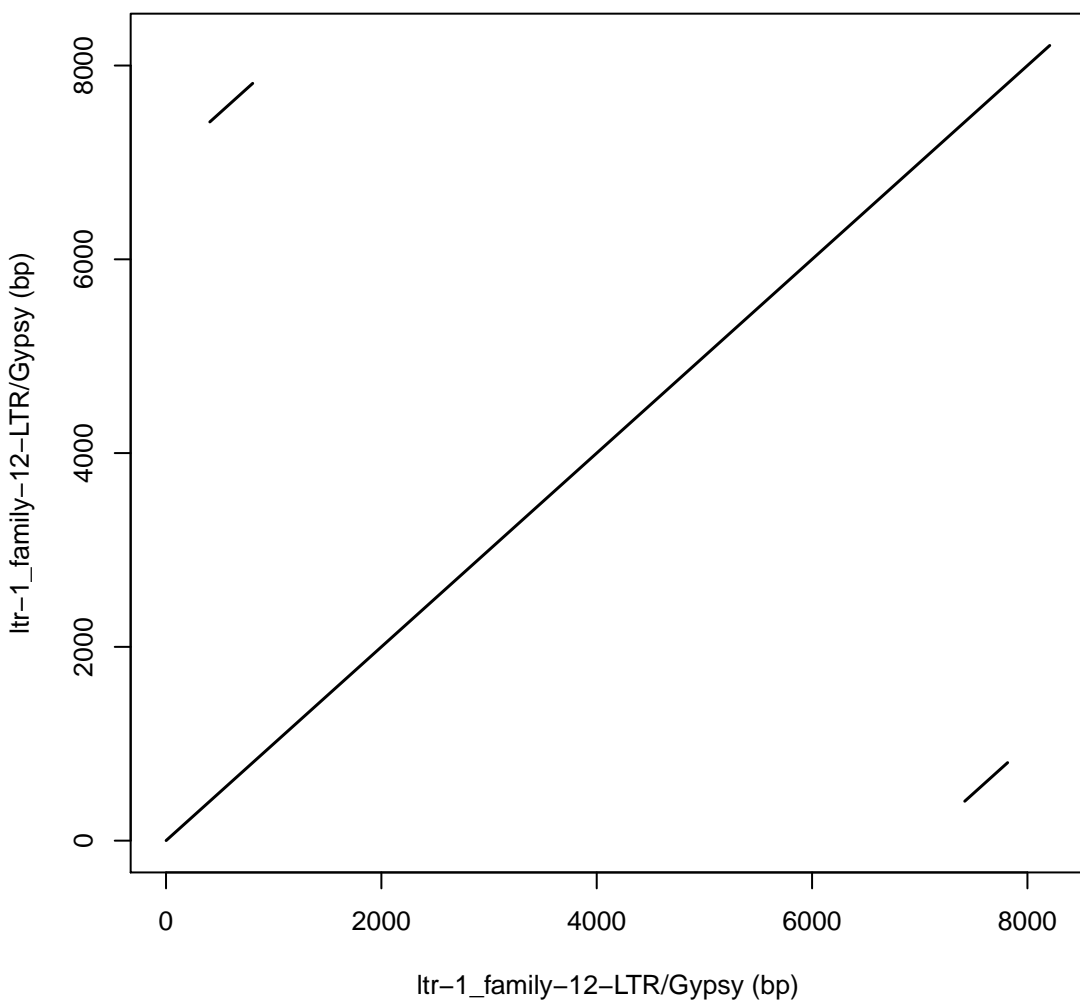
TE: ltr-1_family-12-LTR/Gypsy
consensus size: 8207bp; fragments: 301; full length: 26 (>=7386.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

