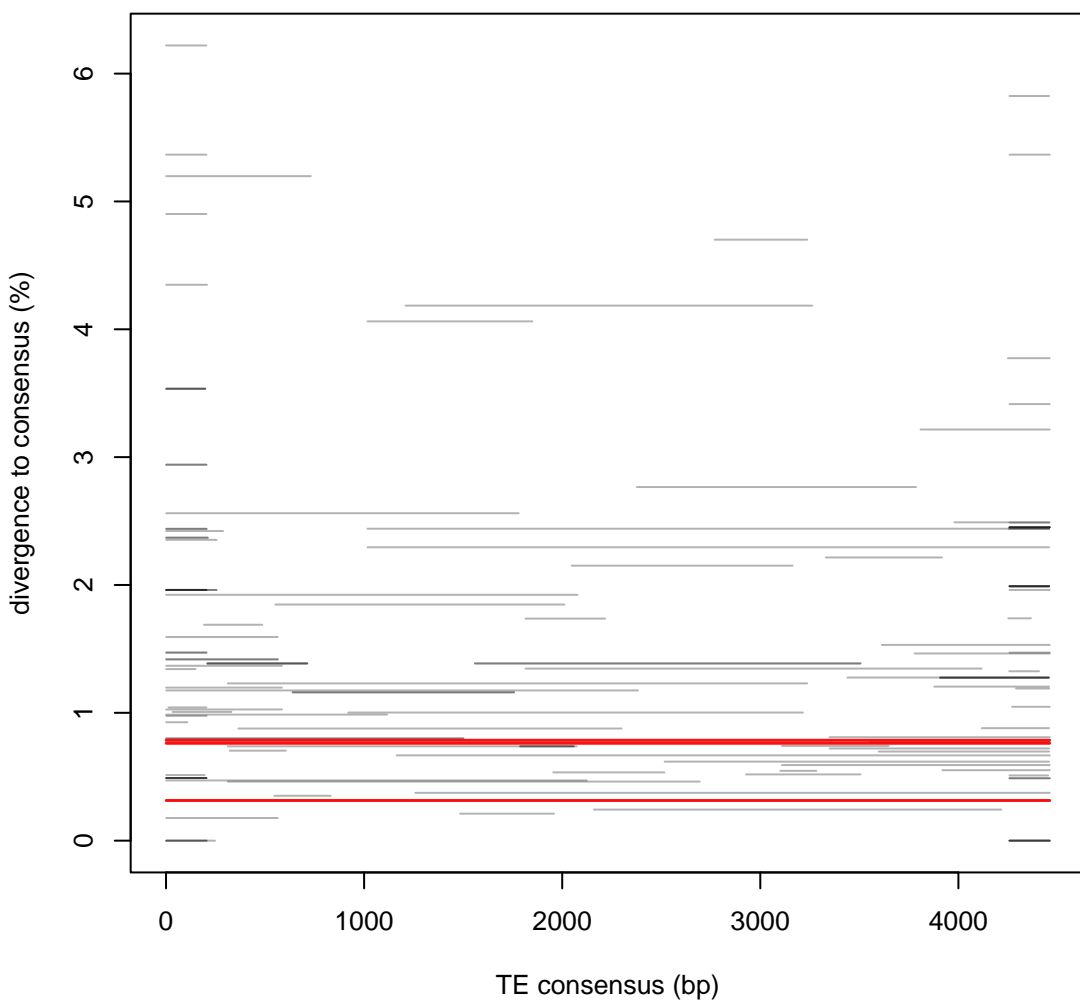
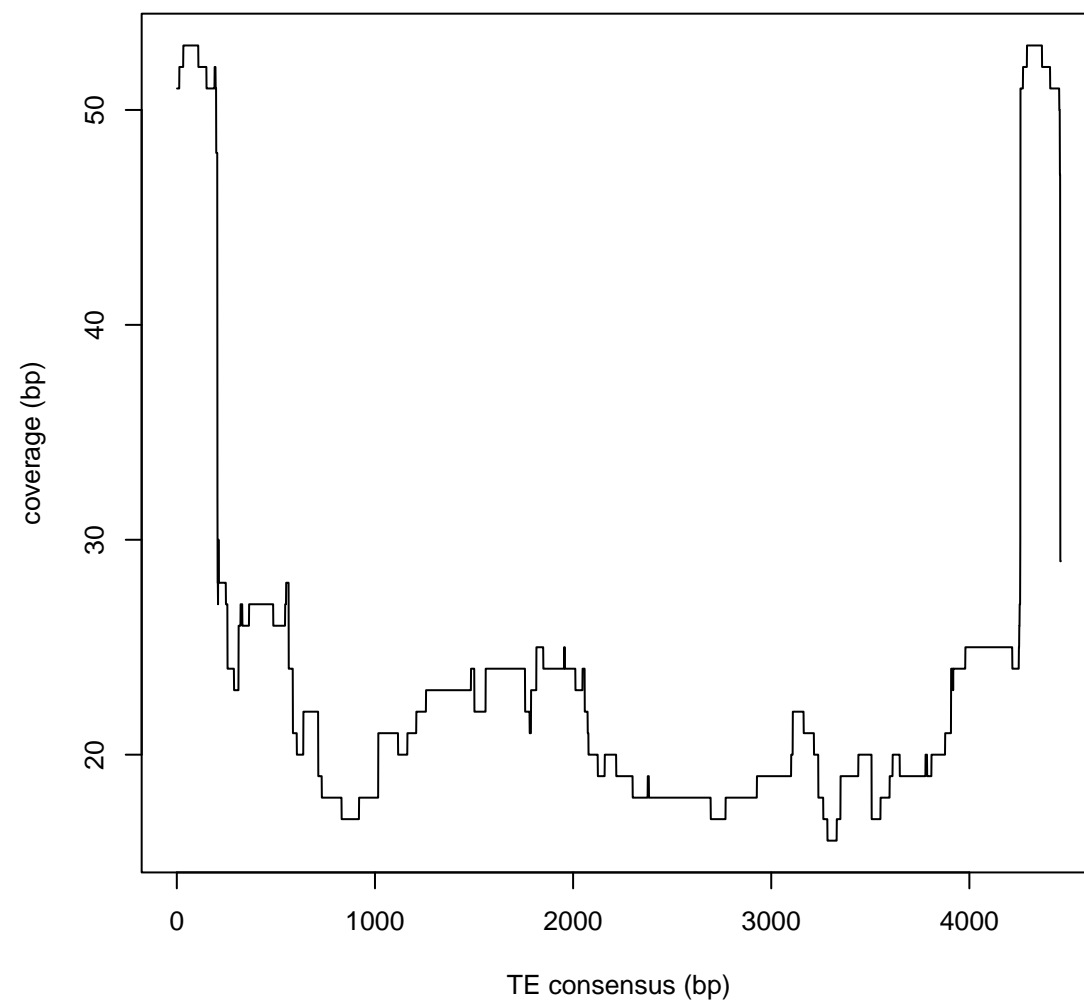


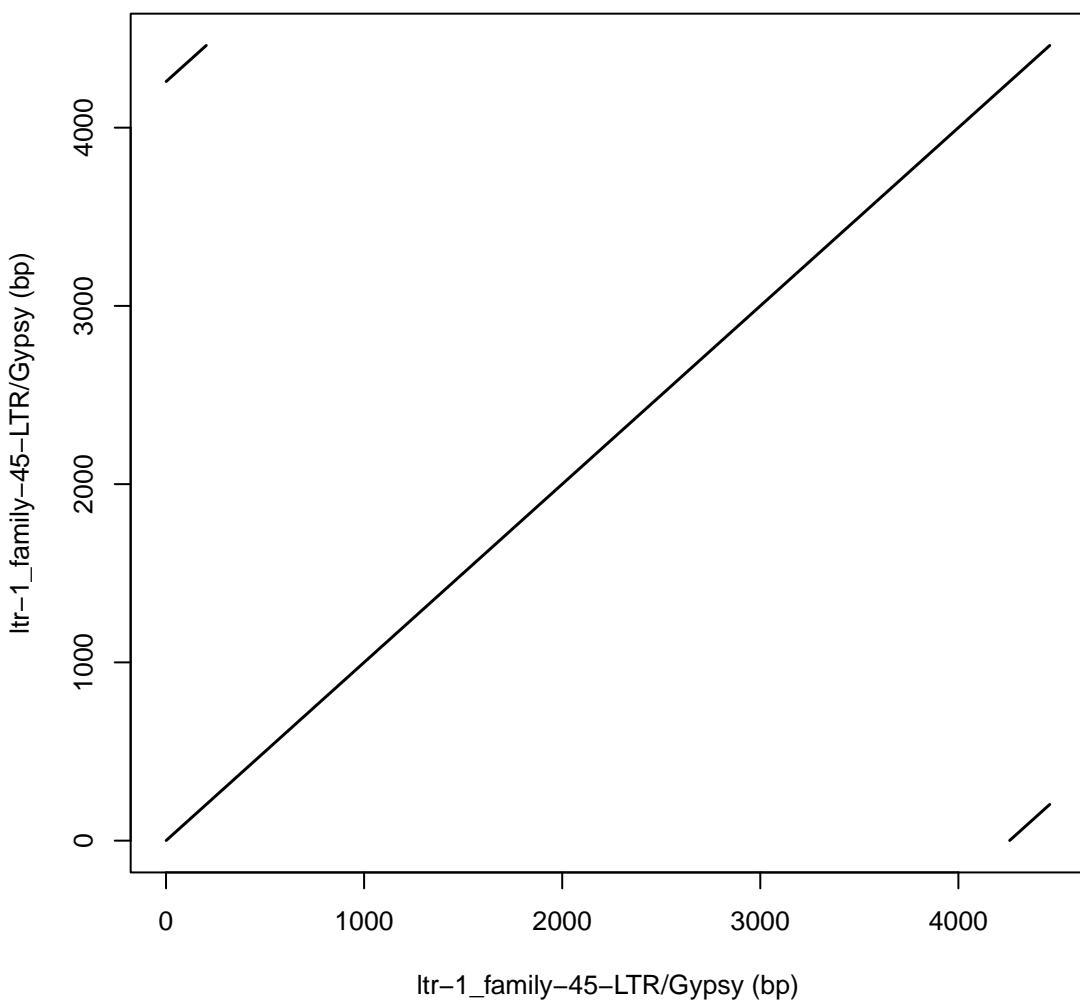
TE: ltr-1_family-45-LTR/Gypsy
consensus size: 4461bp; fragments: 135; full length: 3 (>=4014.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

