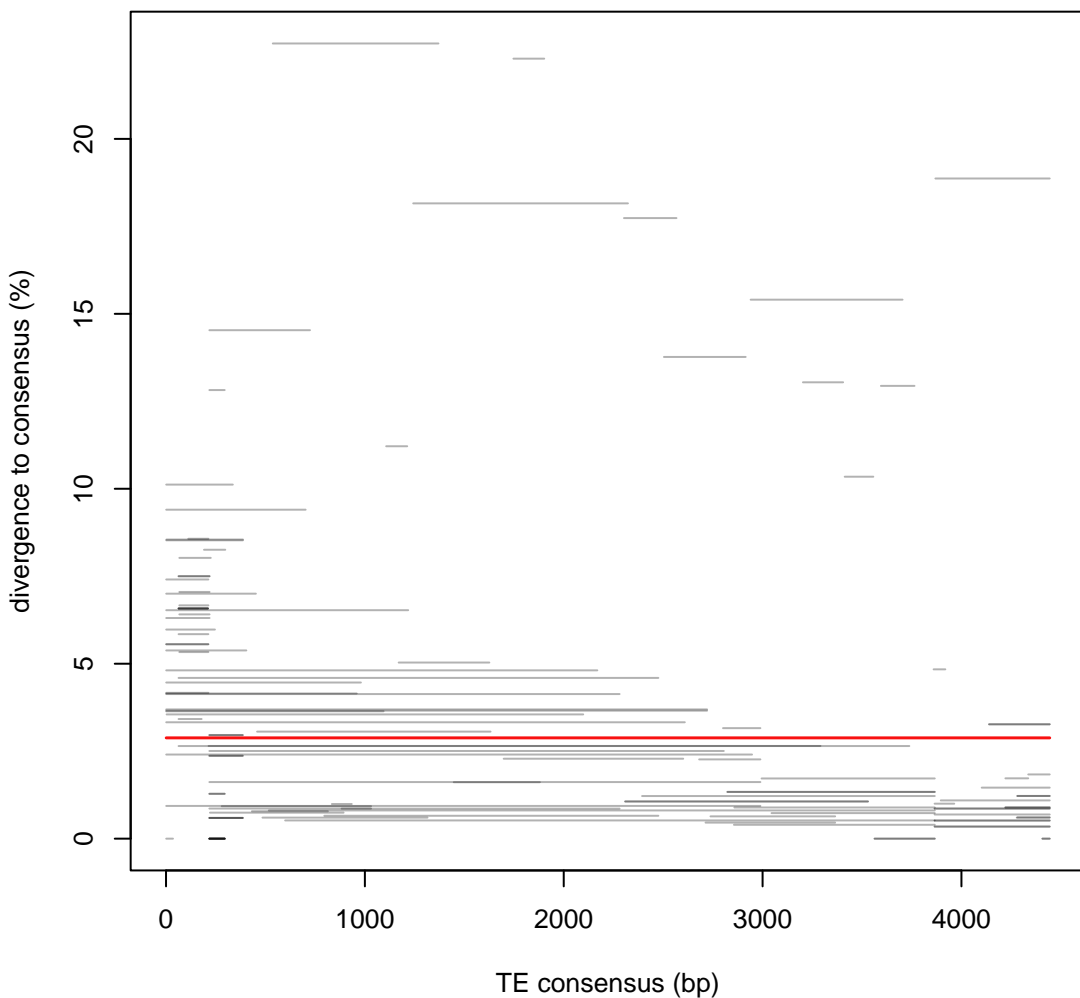
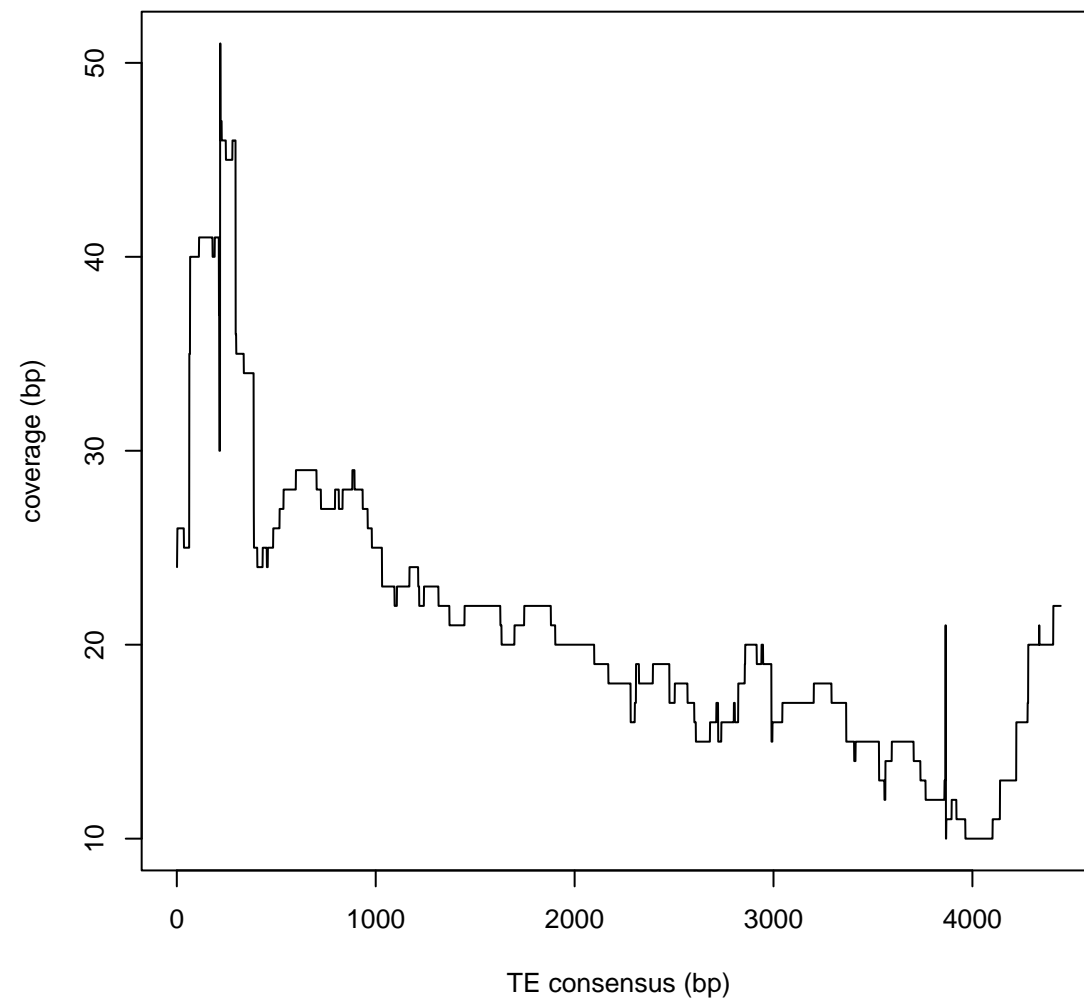


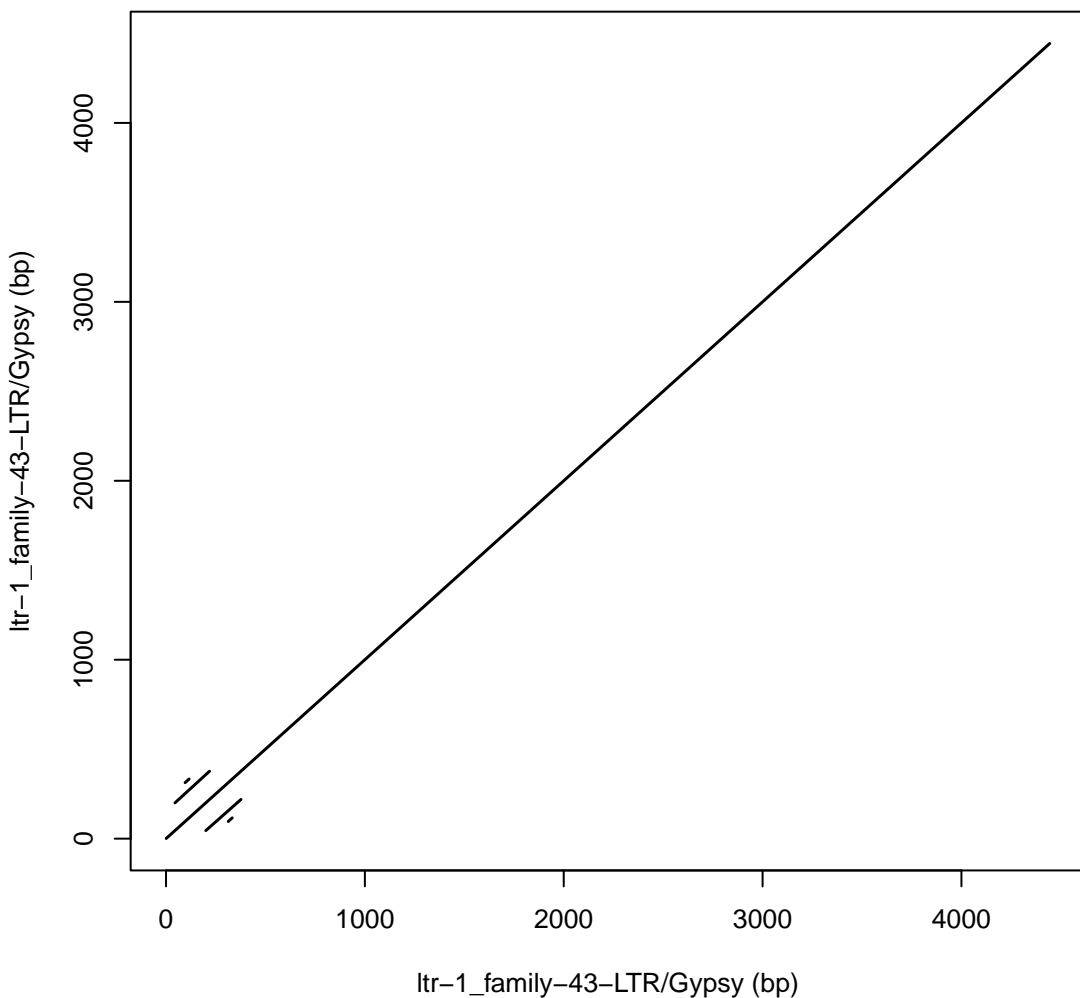
TE: ltr-1_family-43-LTR/Gypsy
consensus size: 4444bp; fragments: 127; full length: 1 (>=3999.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

