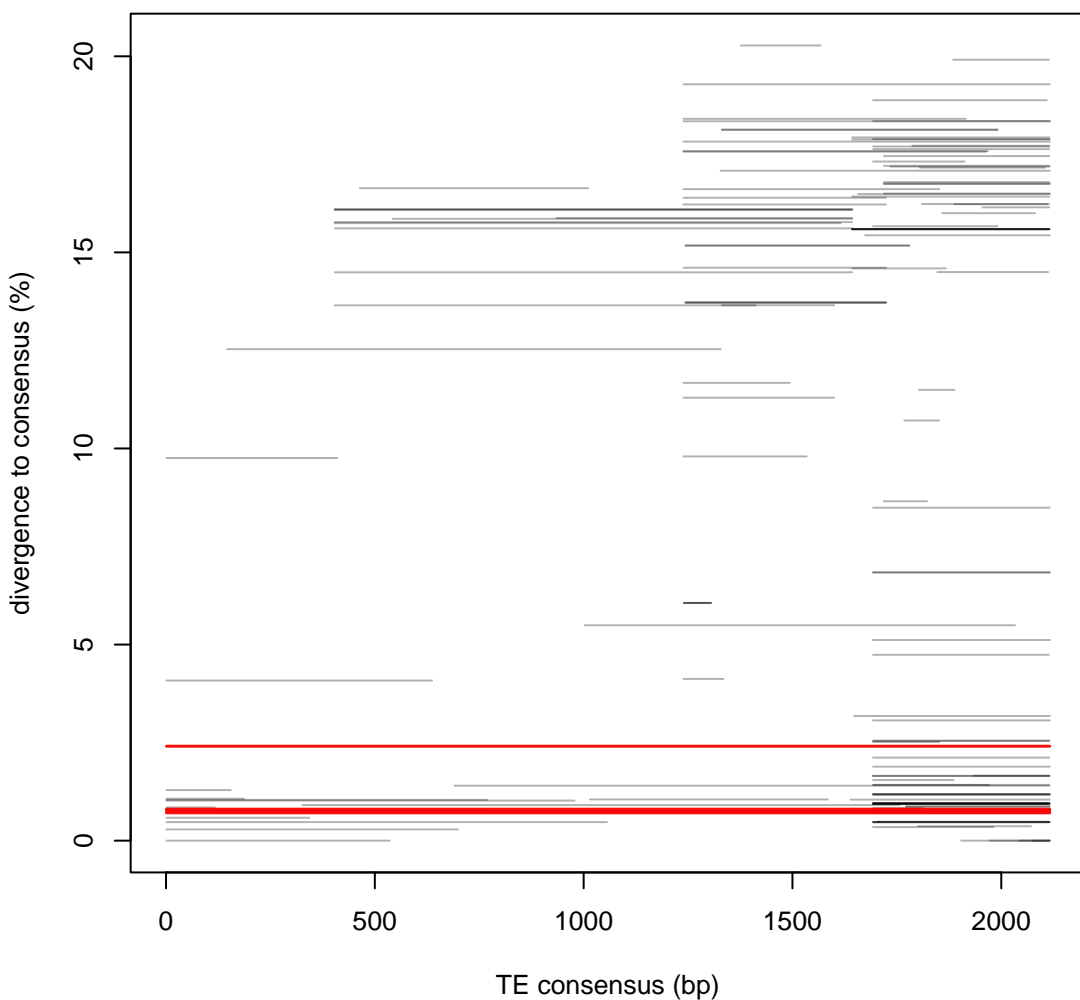
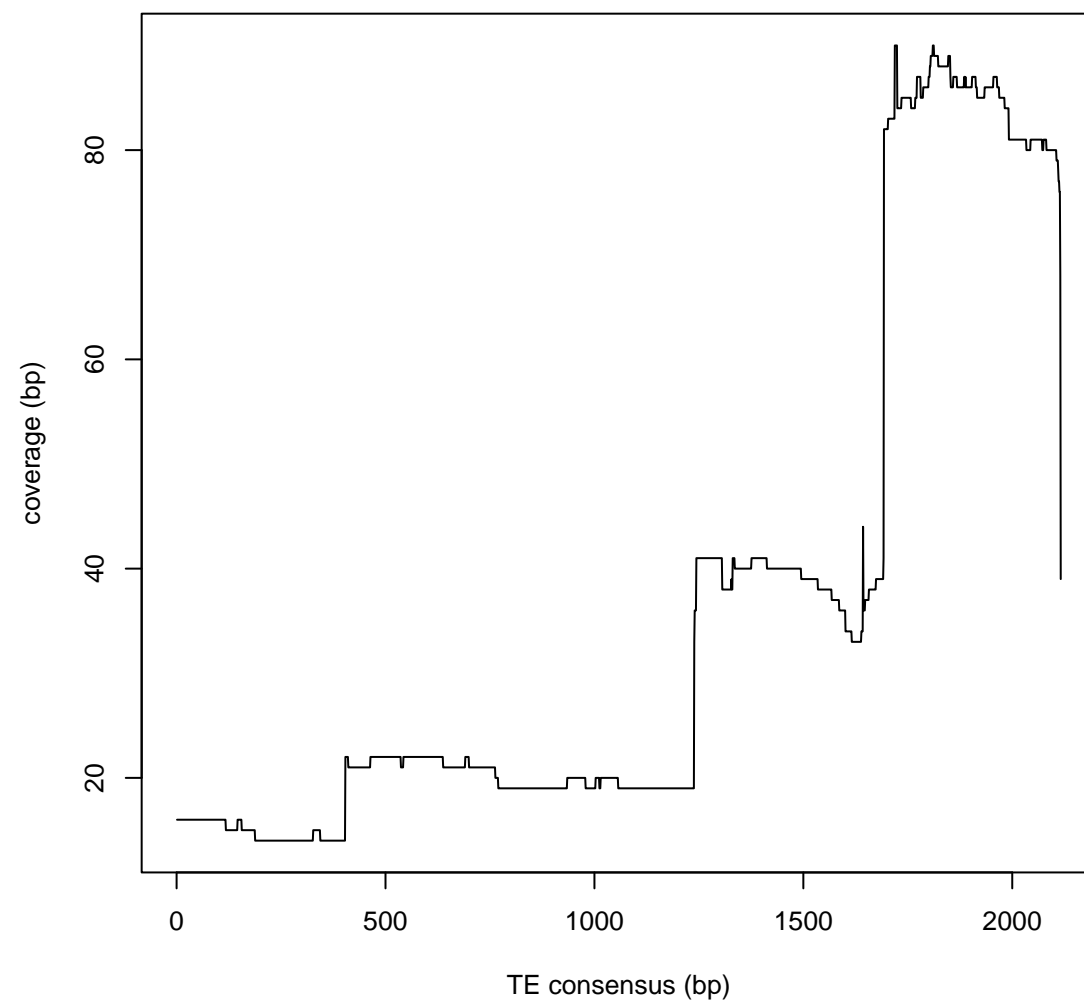


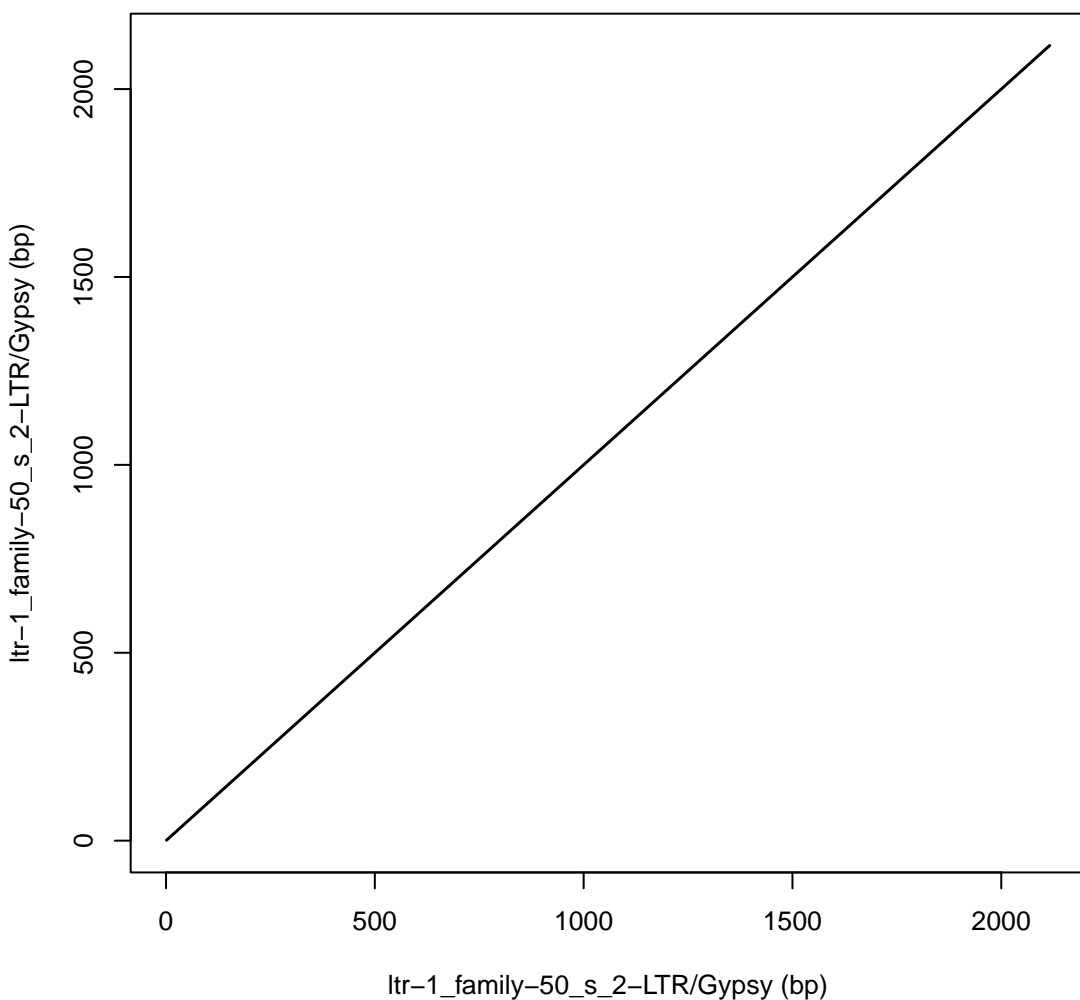
TE: ltr-1\_family-50\_s\_2-LTR/Gypsy  
consensus size: 2116bp; fragments: 143; full length: 4 (>=1904.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

