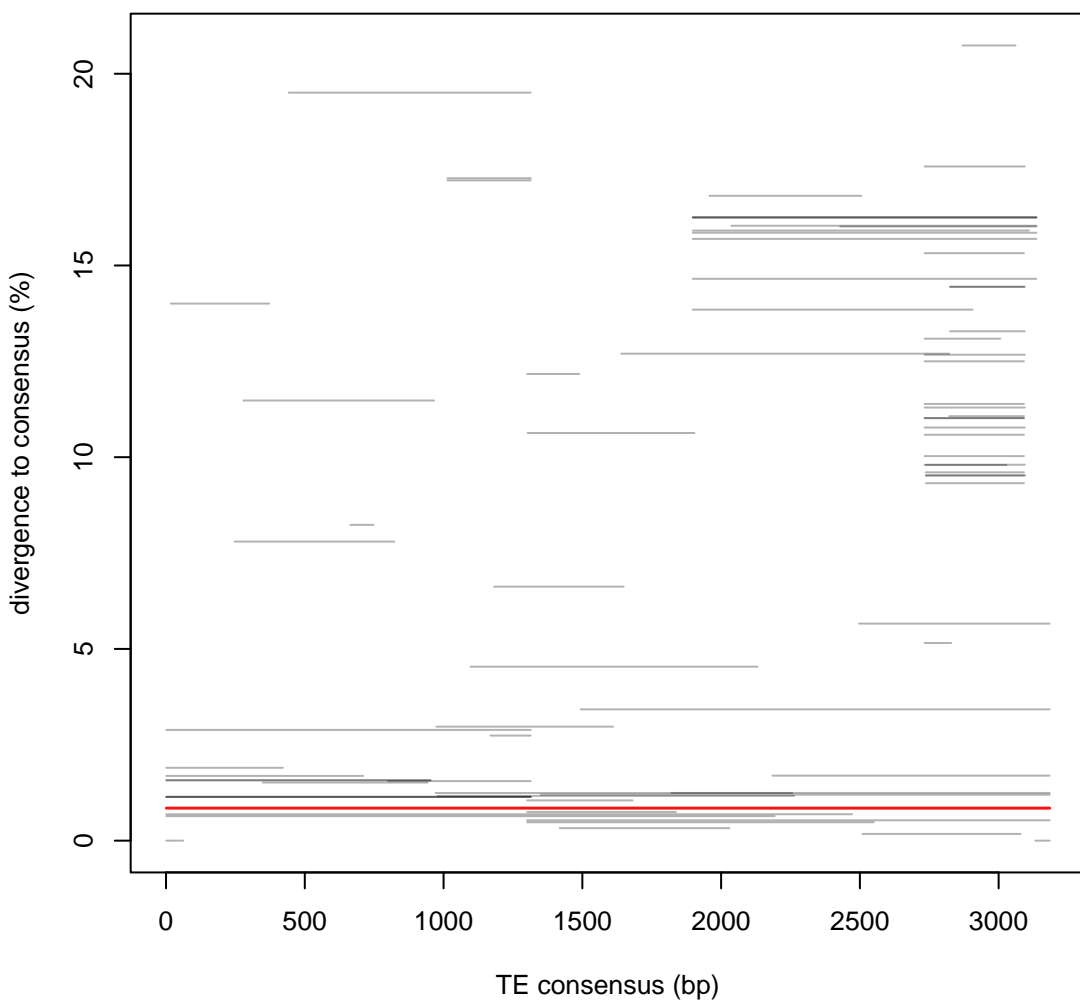
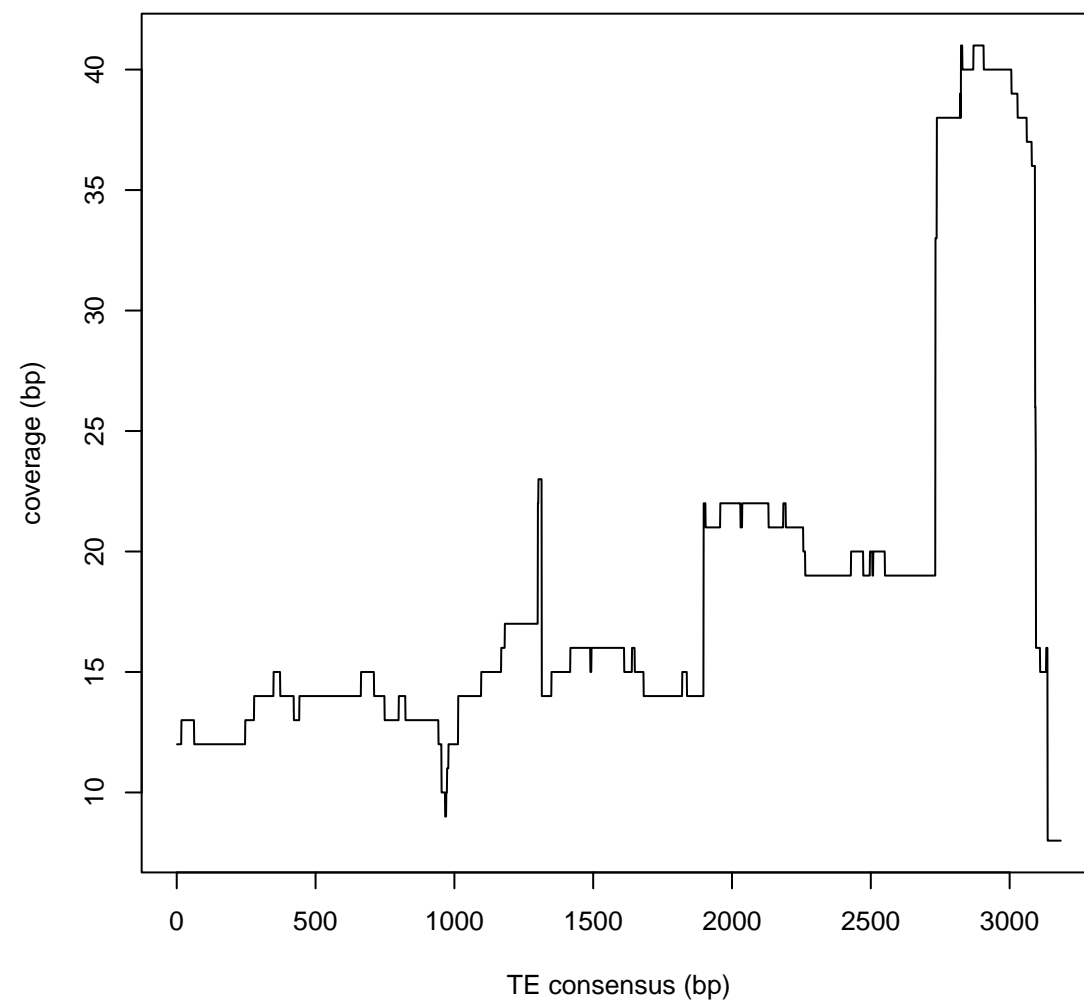


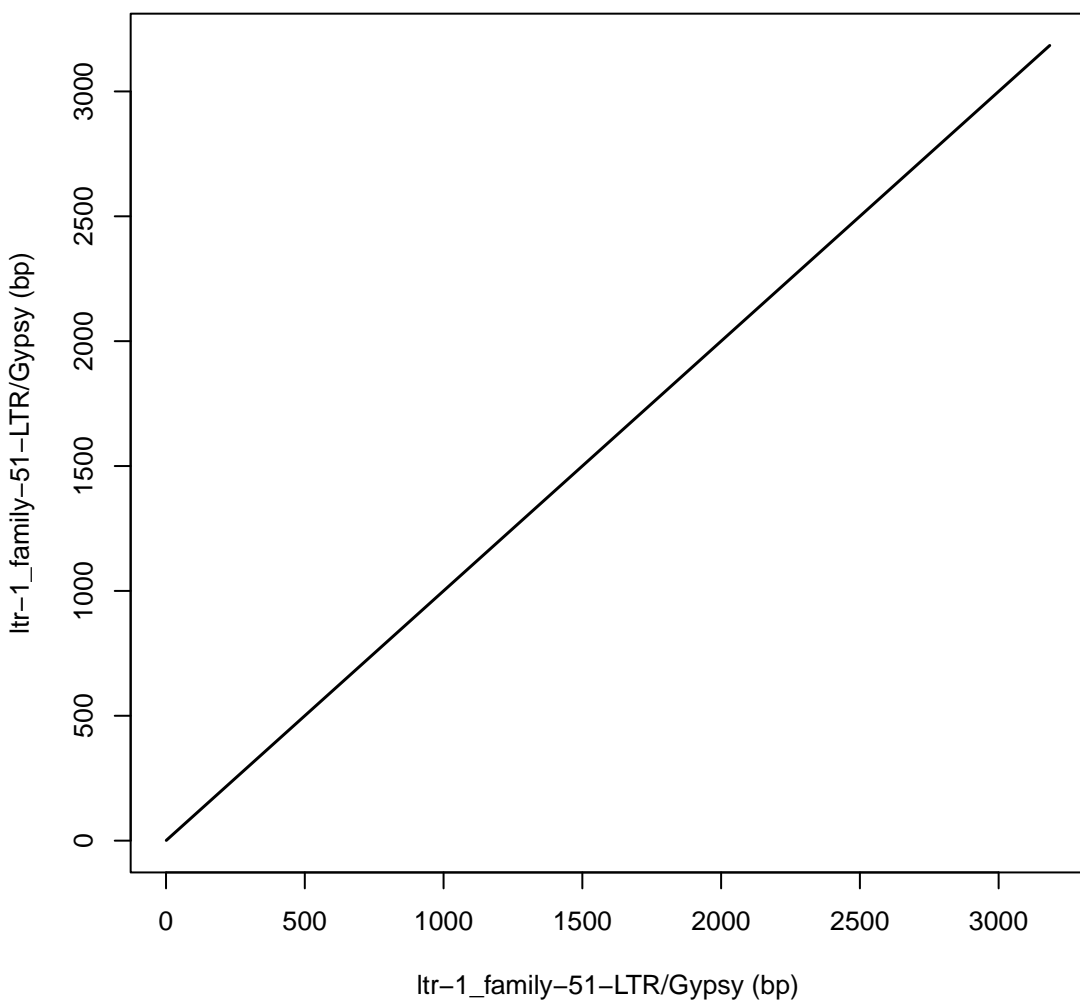
TE: ltr-1_family-51-LTR/Gypsy
consensus size: 3184bp; fragments: 77; full length: 1 (>=2865.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

