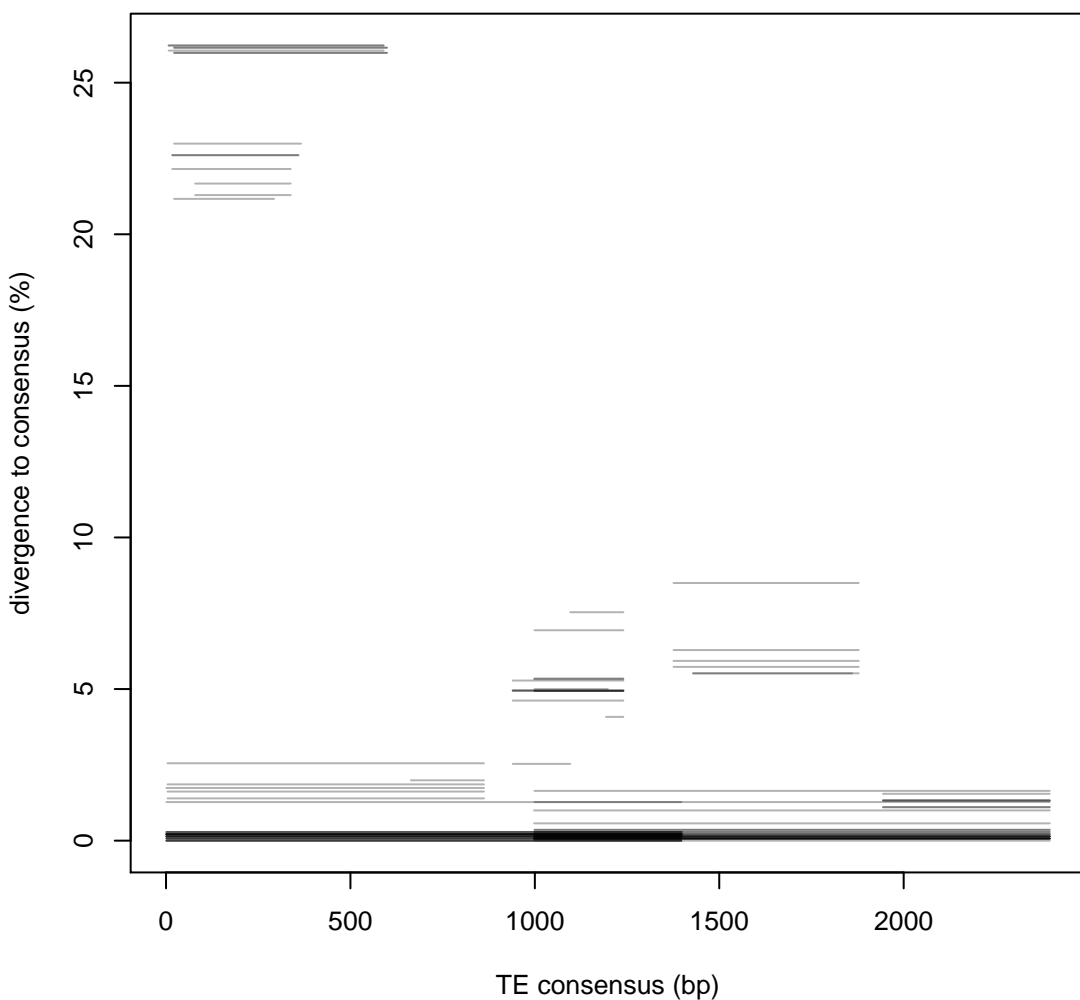
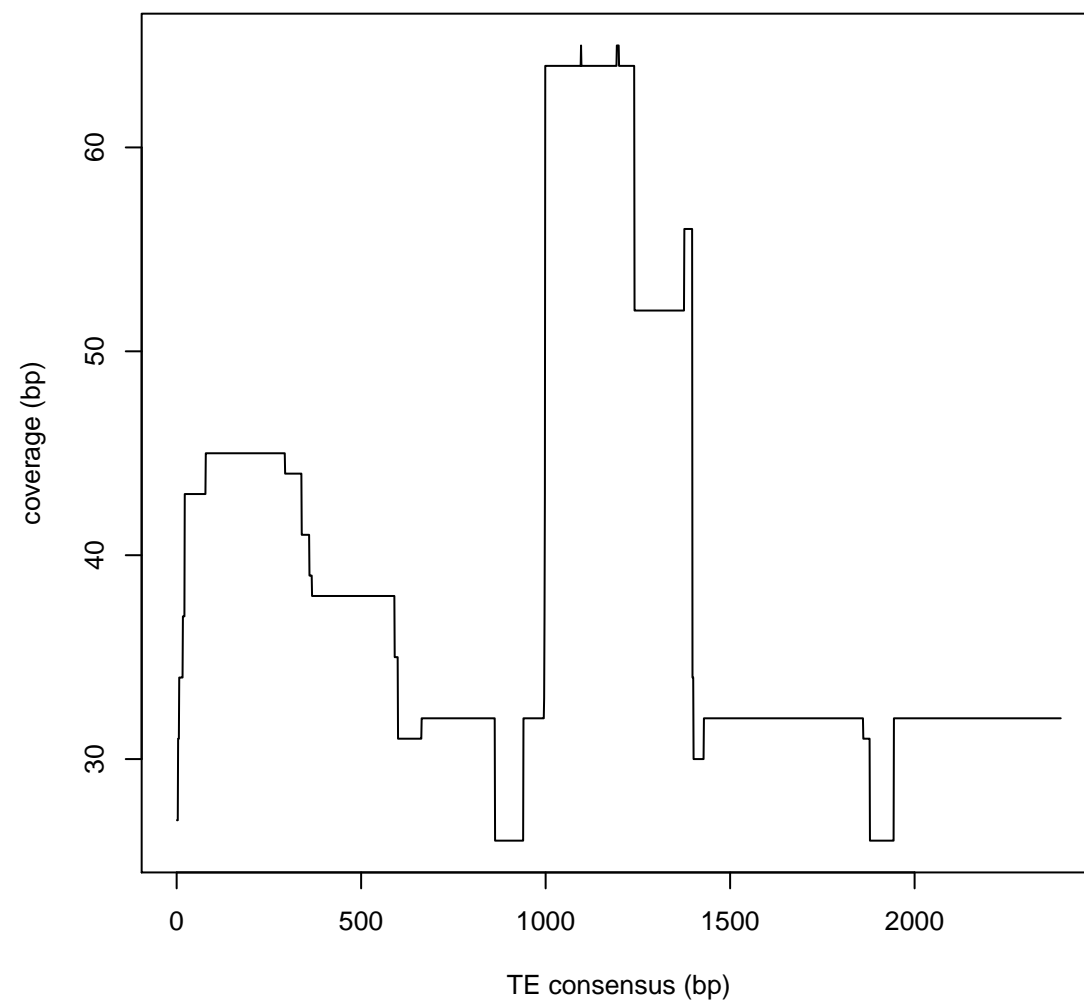


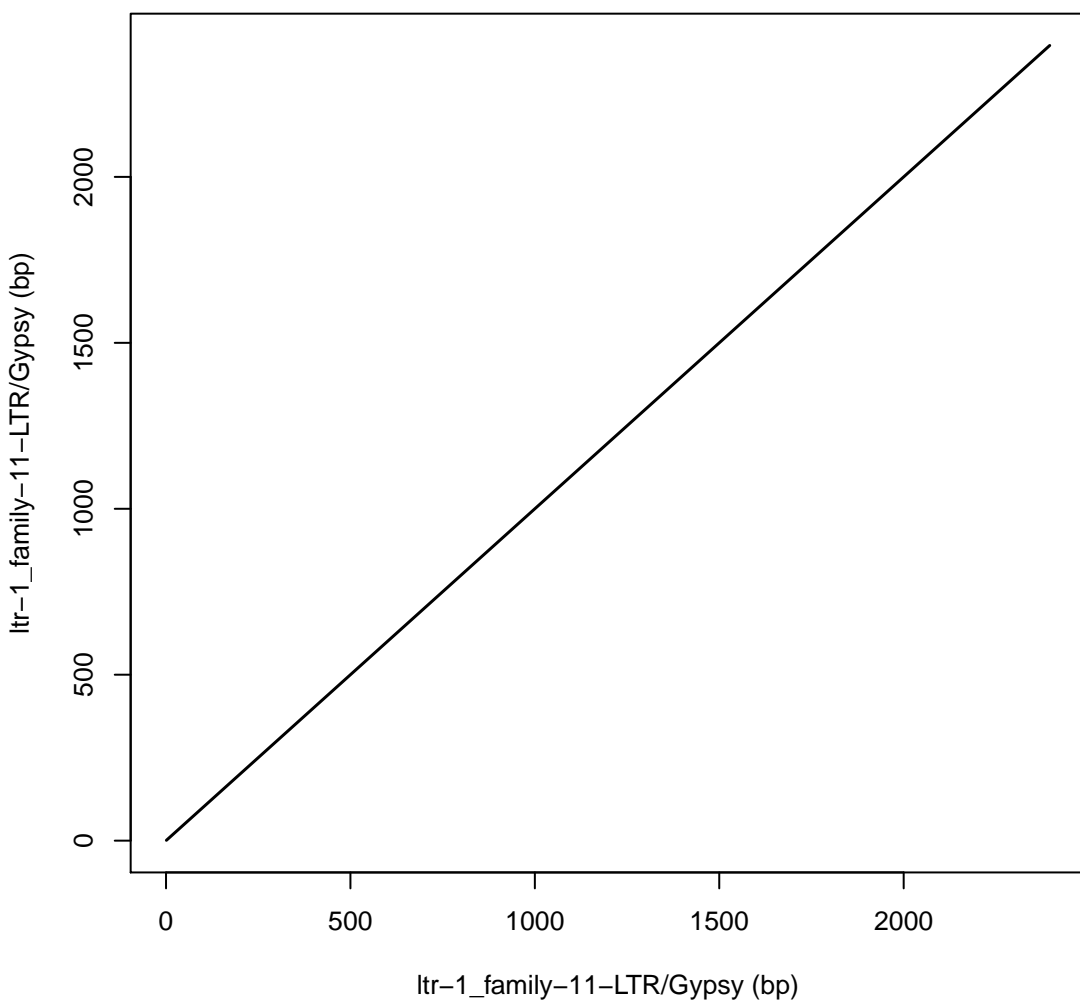
TE: ltr-1\_family-11-LTR/Gypsy  
consensus size: 2396bp; fragments: 98; full length: 0 (>=2156.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

