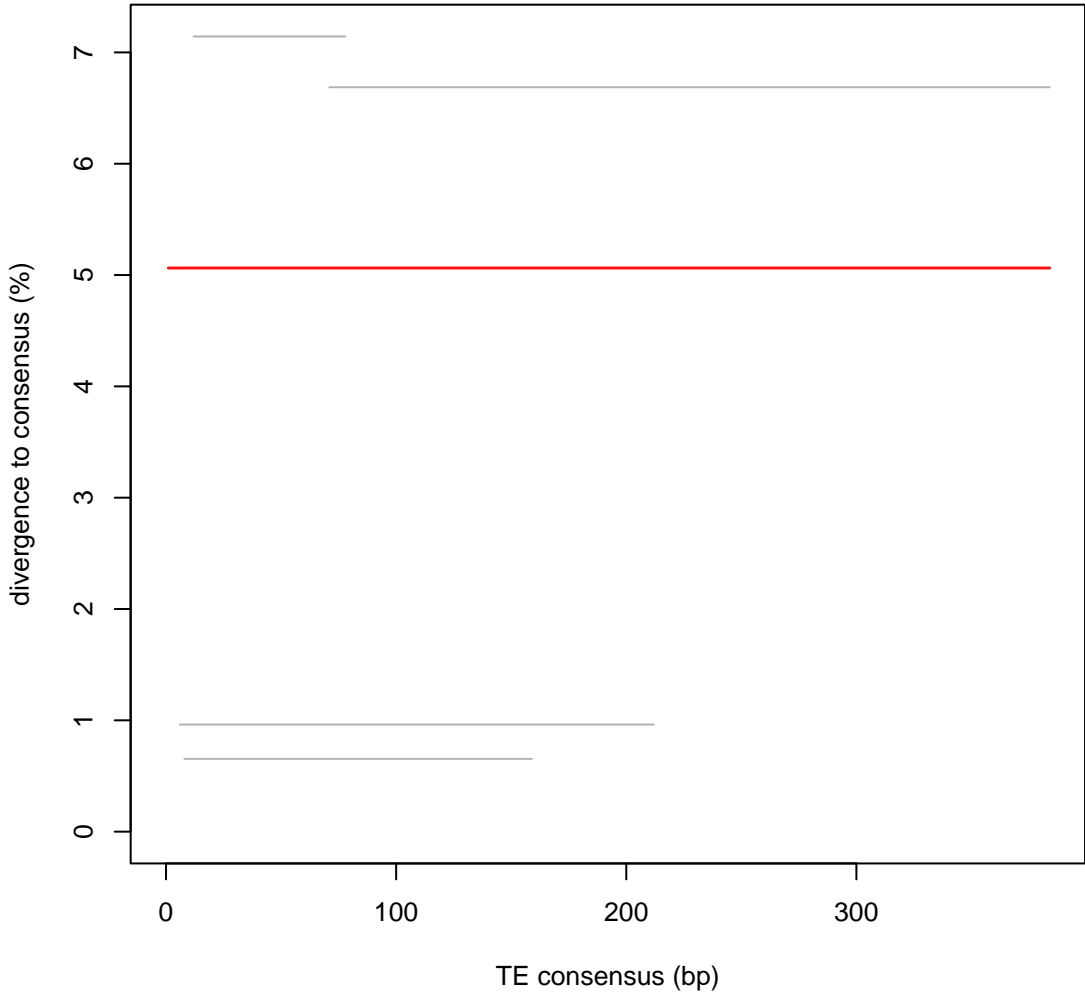
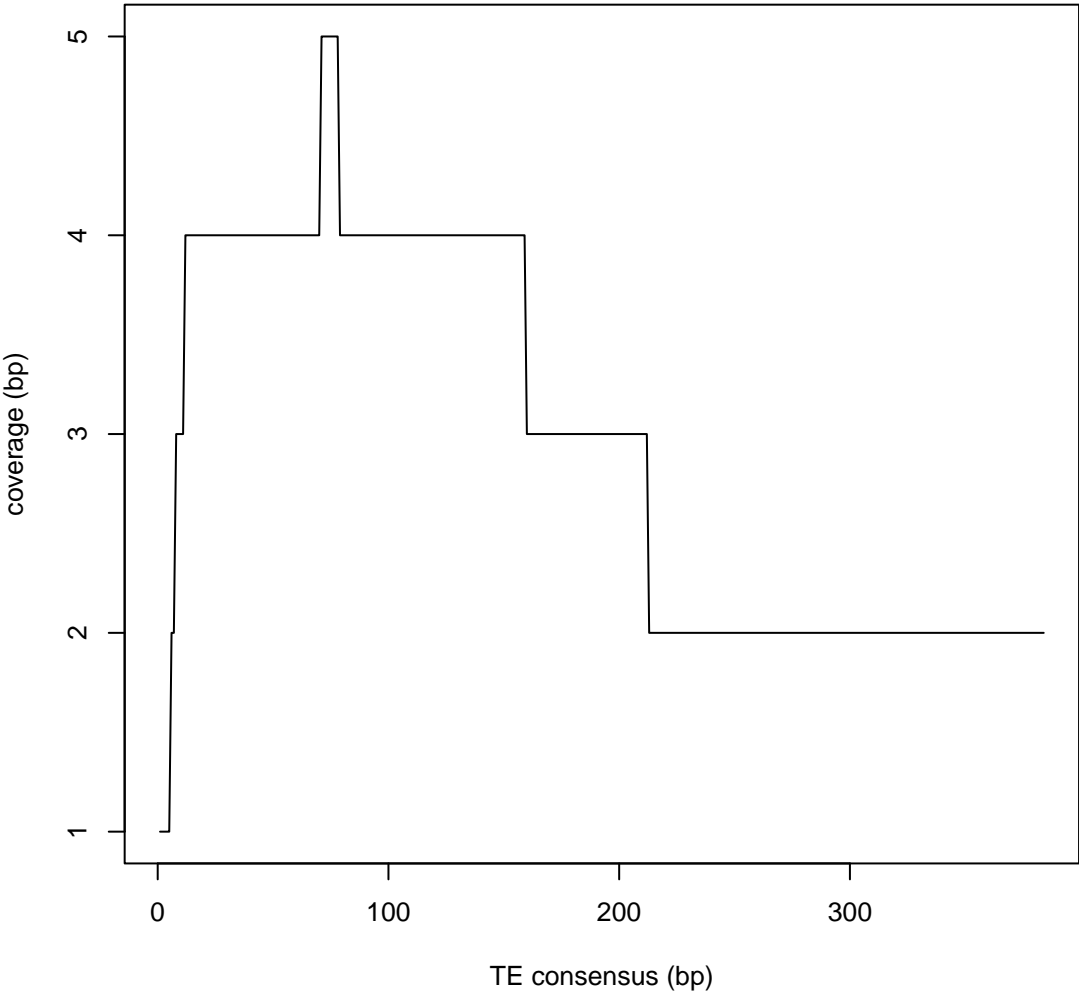


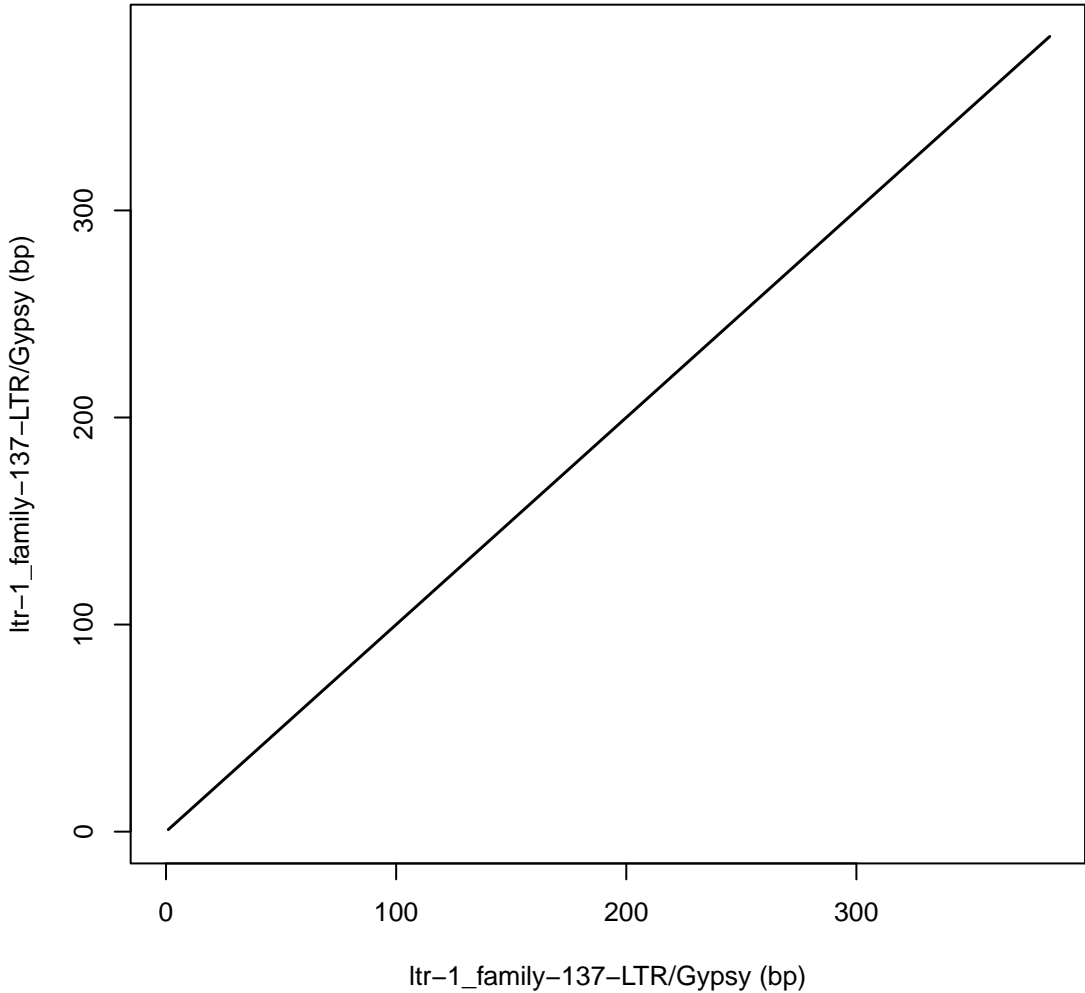
TE: ltr-1\_family-137-LTR/Gypsy  
consensus size: 384bp; fragments: 5; full length: 1 (>=345.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

