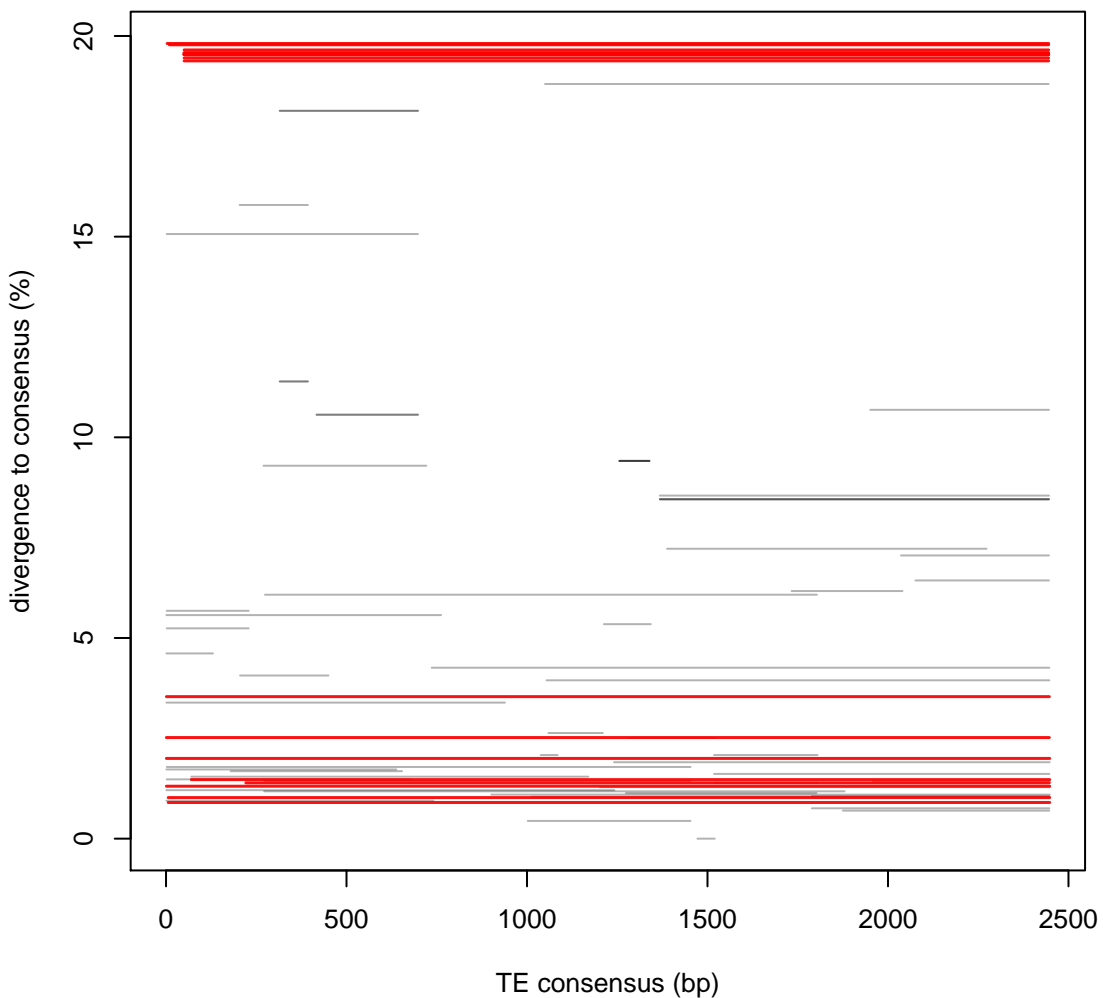
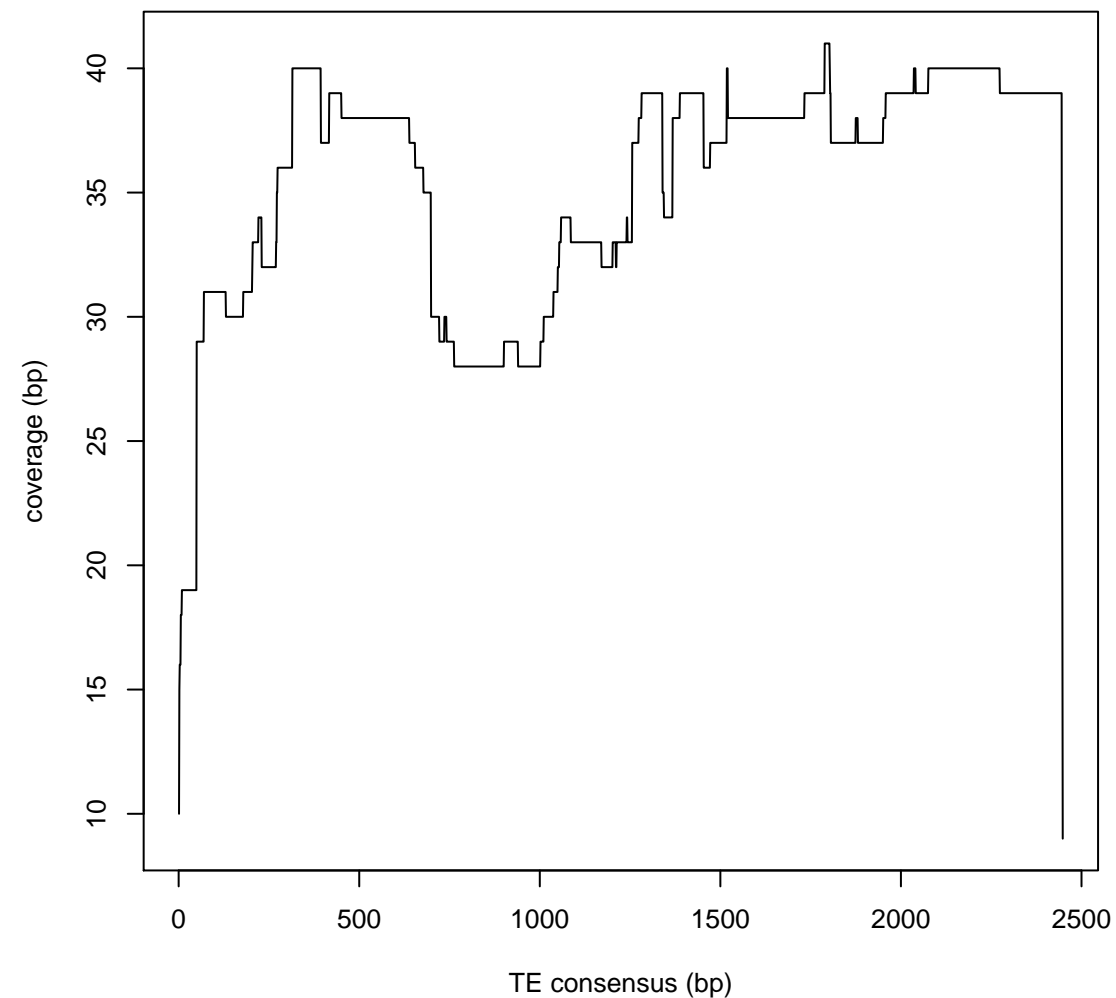


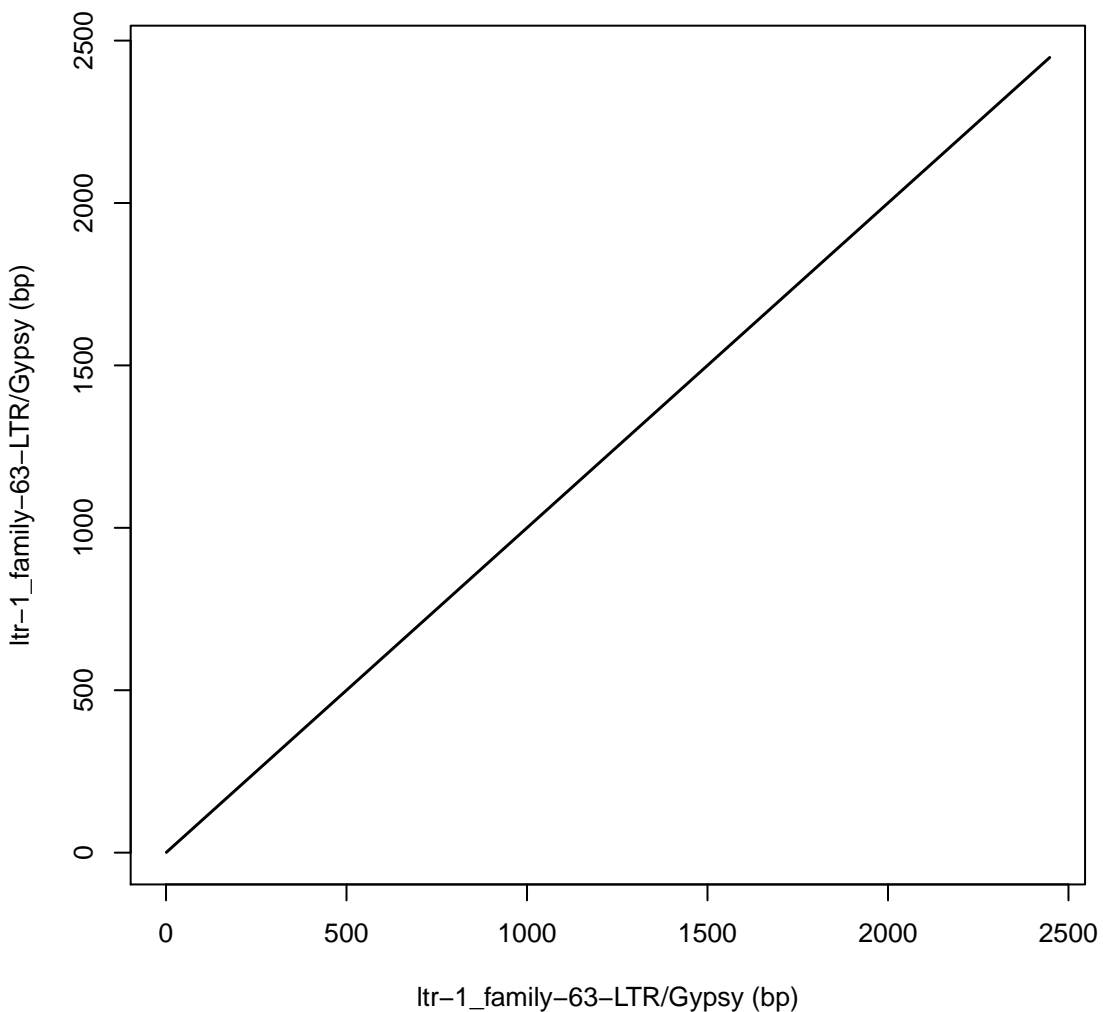
TE: ltr-1\_family-63-LTR/Gypsy  
consensus size: 2448bp; fragments: 79; full length: 20 ( $\geq 2203.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

