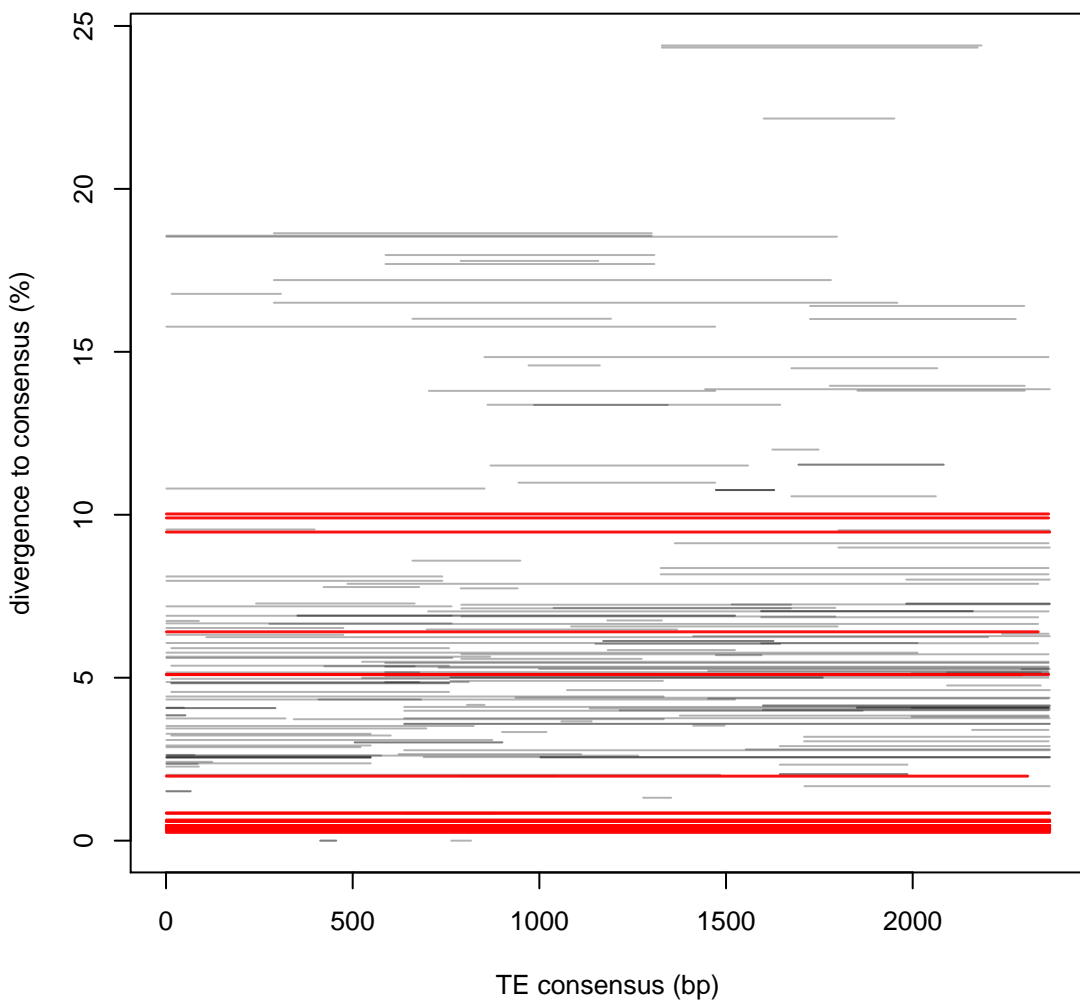
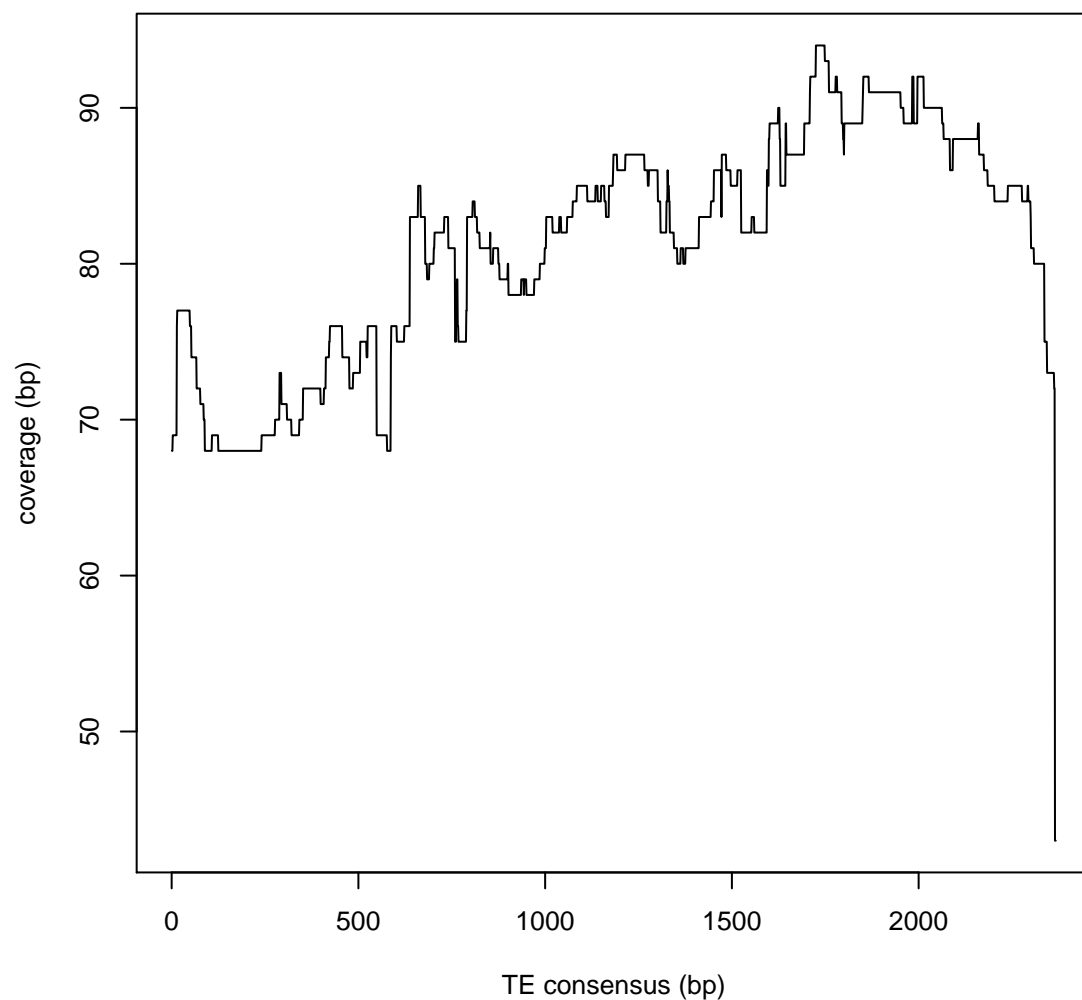


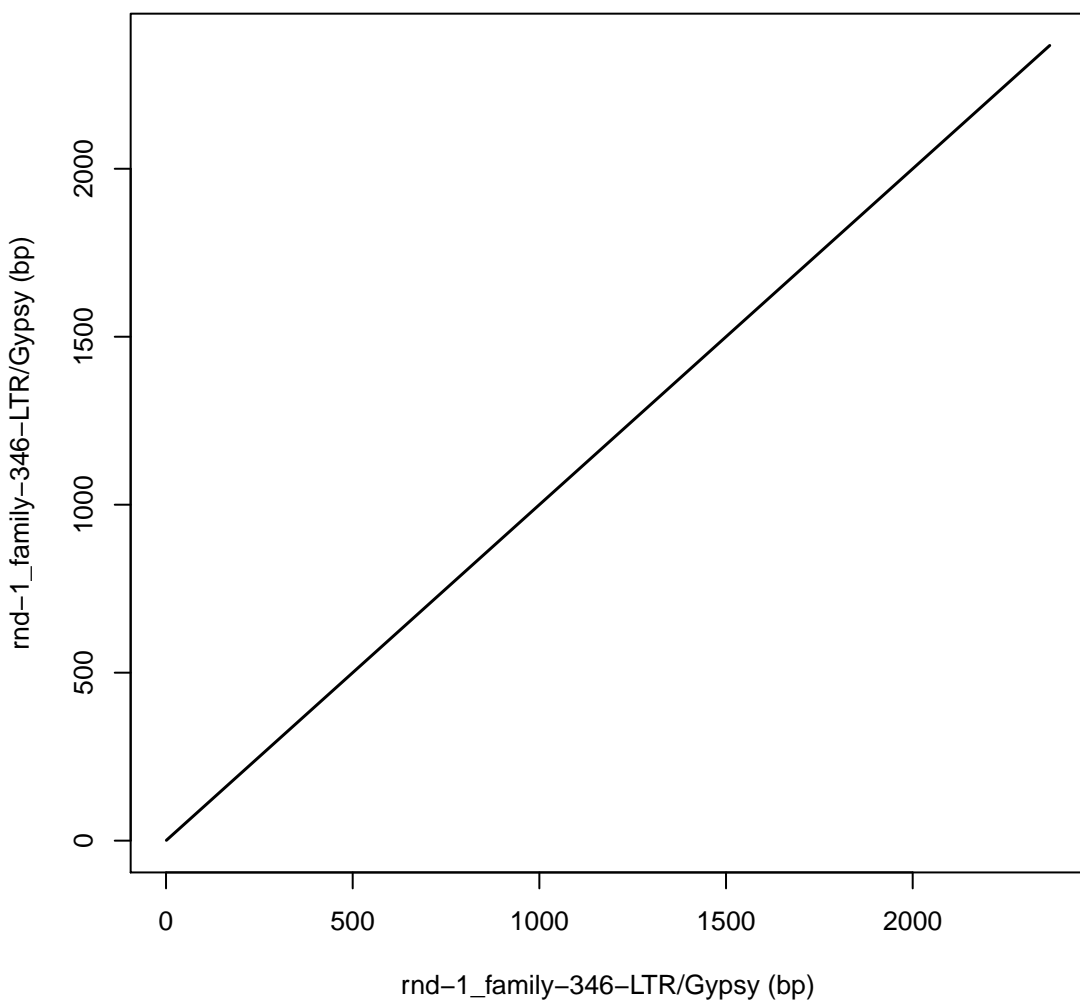
TE: rnd-1\_family-346-LTR/Gypsy  
consensus size: 2367bp; fragments: 211; full length: 22 ( $\geq 2130.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

