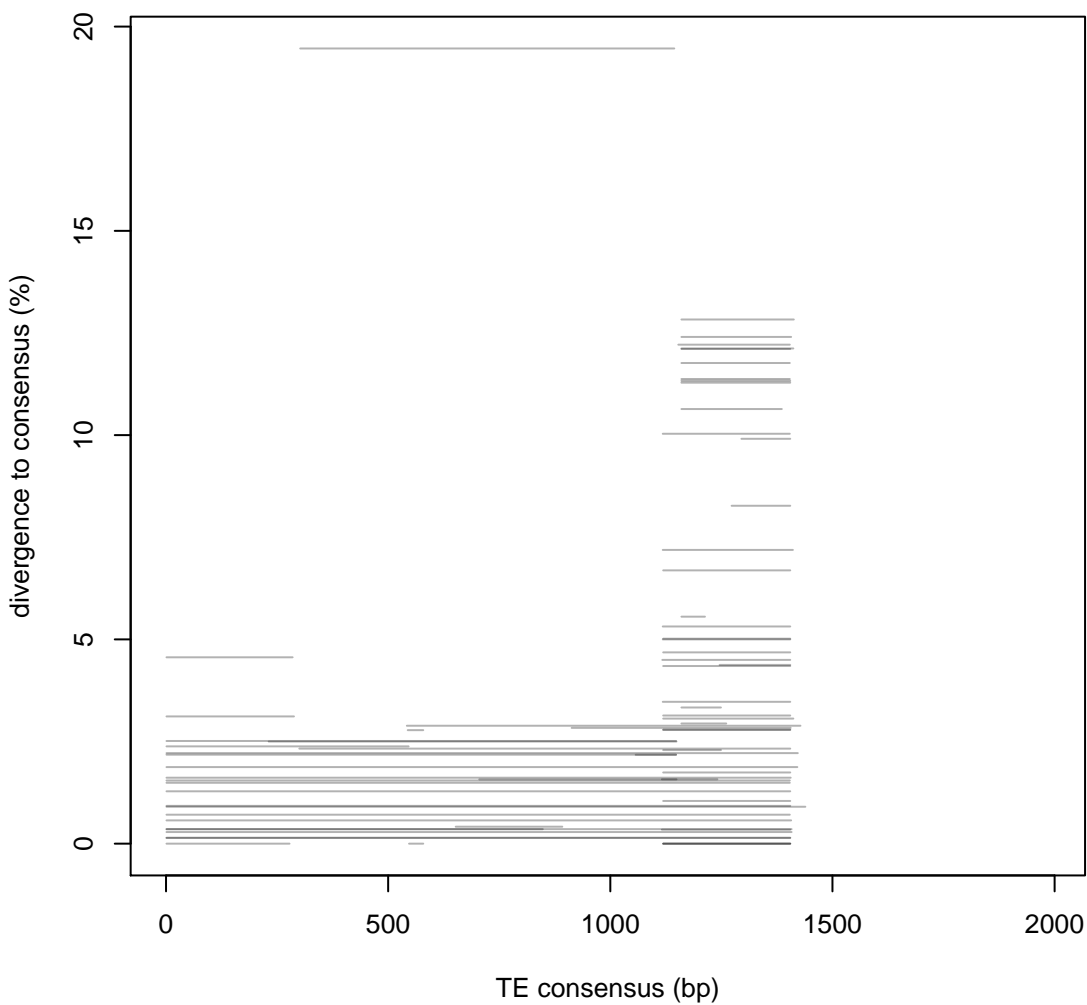
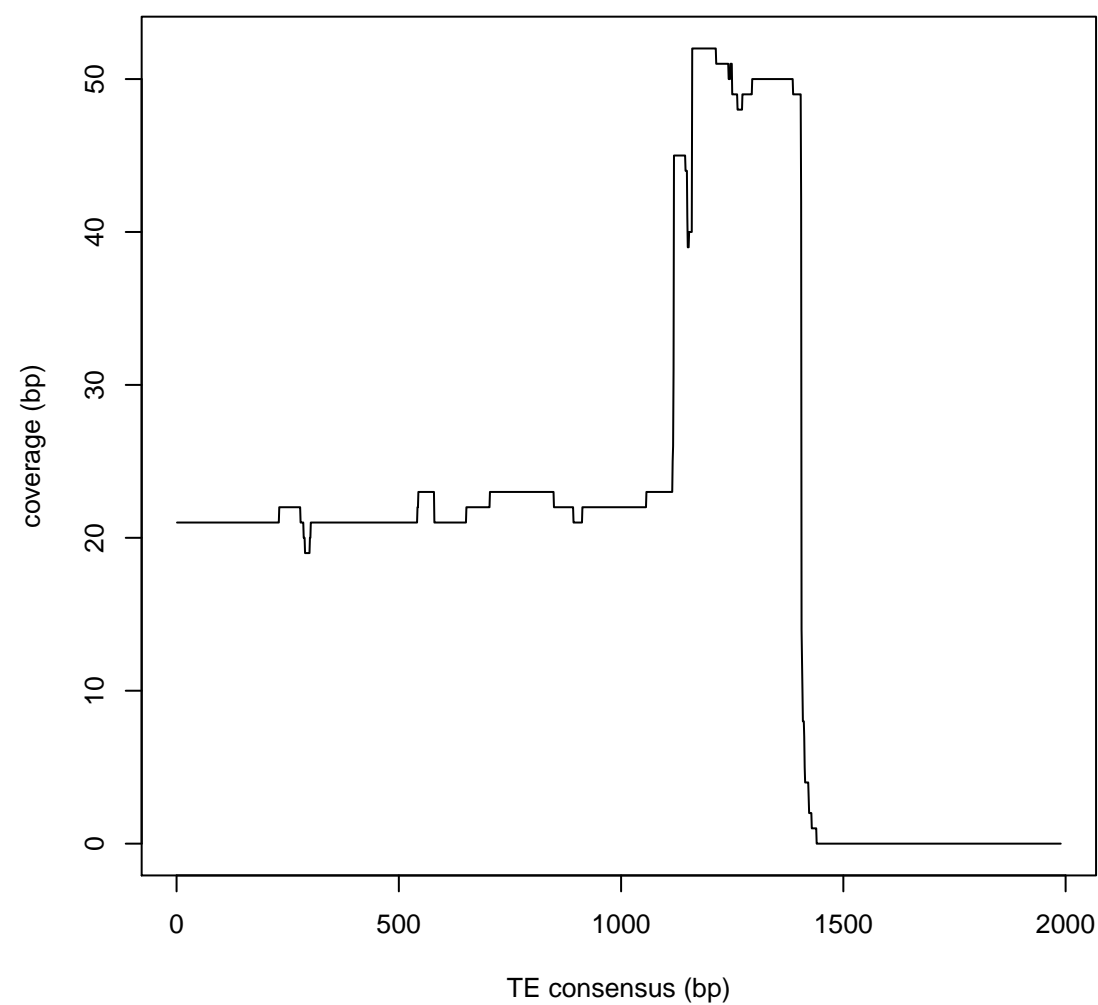


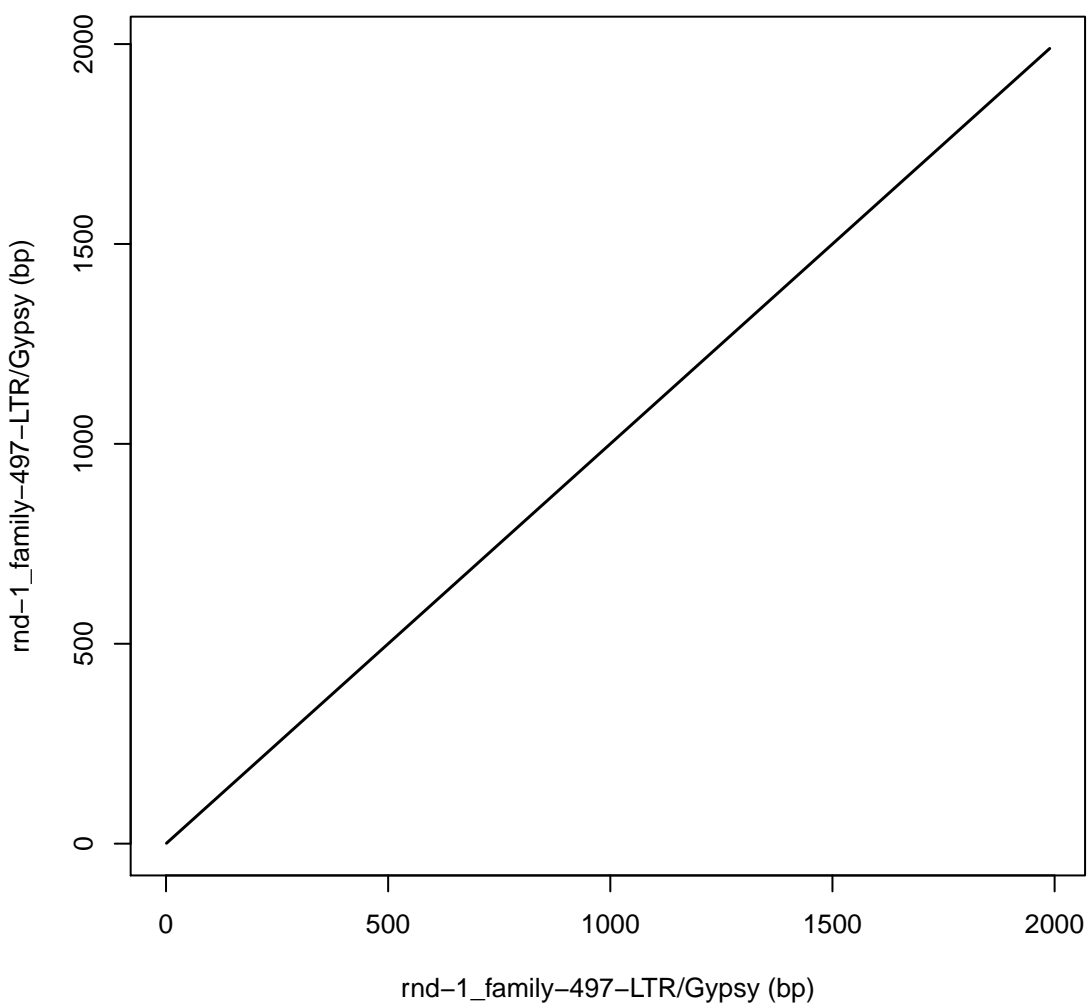
TE: rnd-1\_family-497-LTR/Gypsy  
consensus size: 1989bp; fragments: 69; full length: 0 ( $\geq 1790.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

