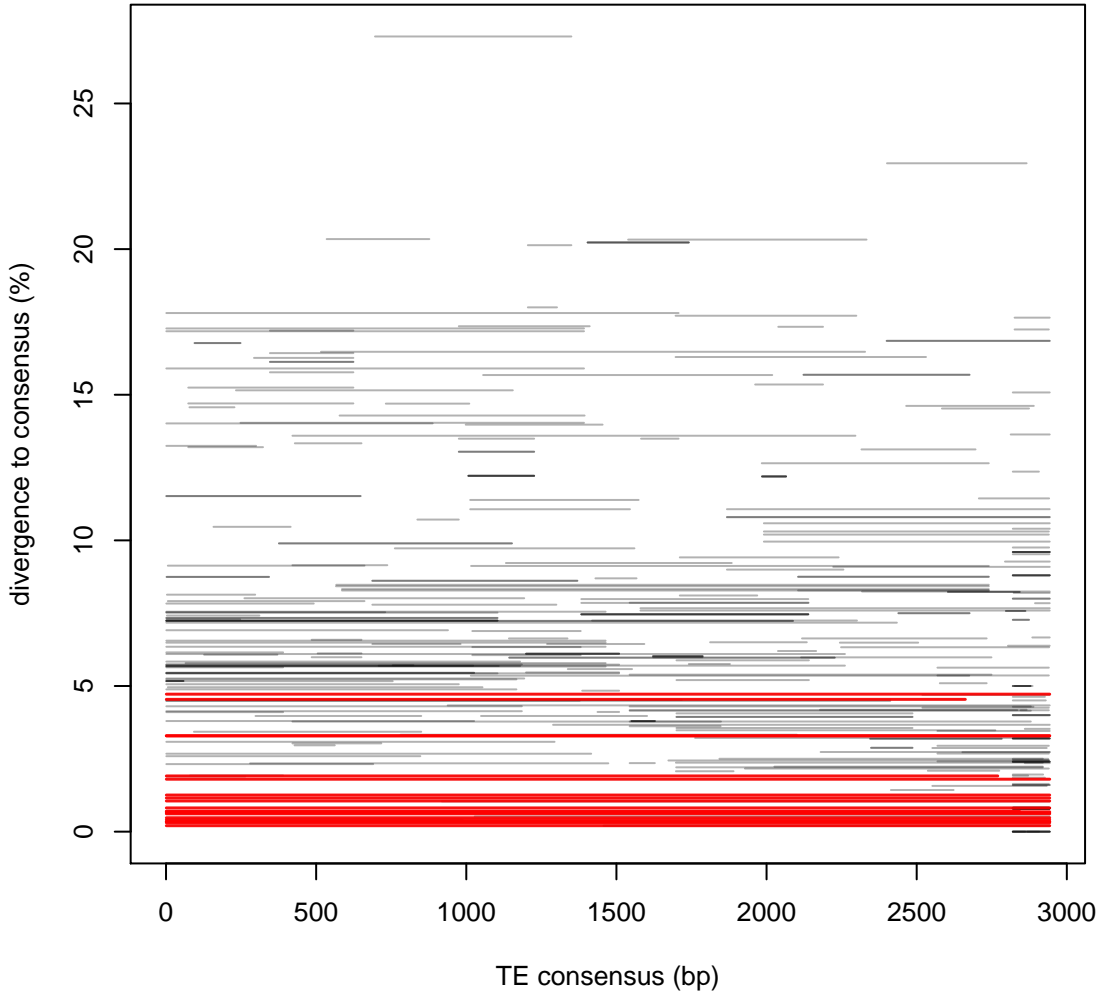
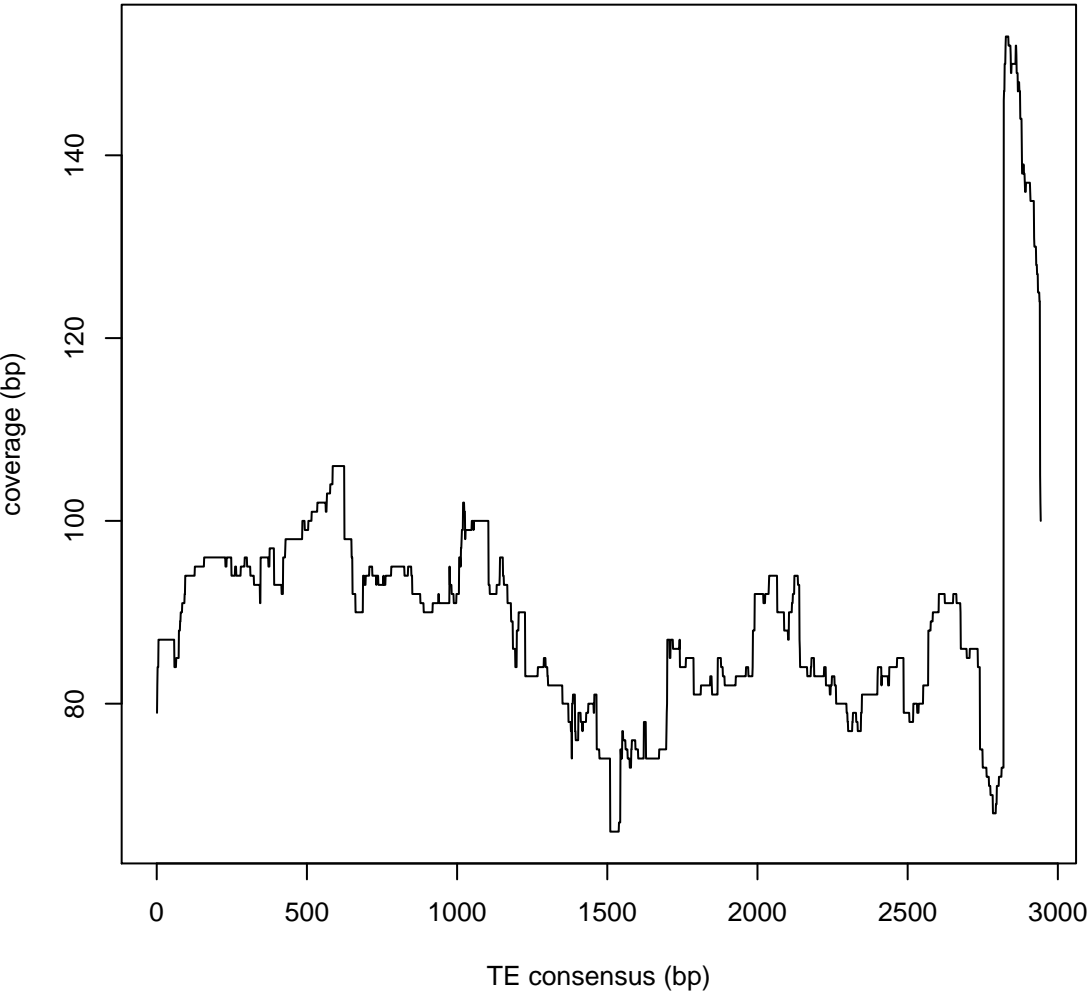


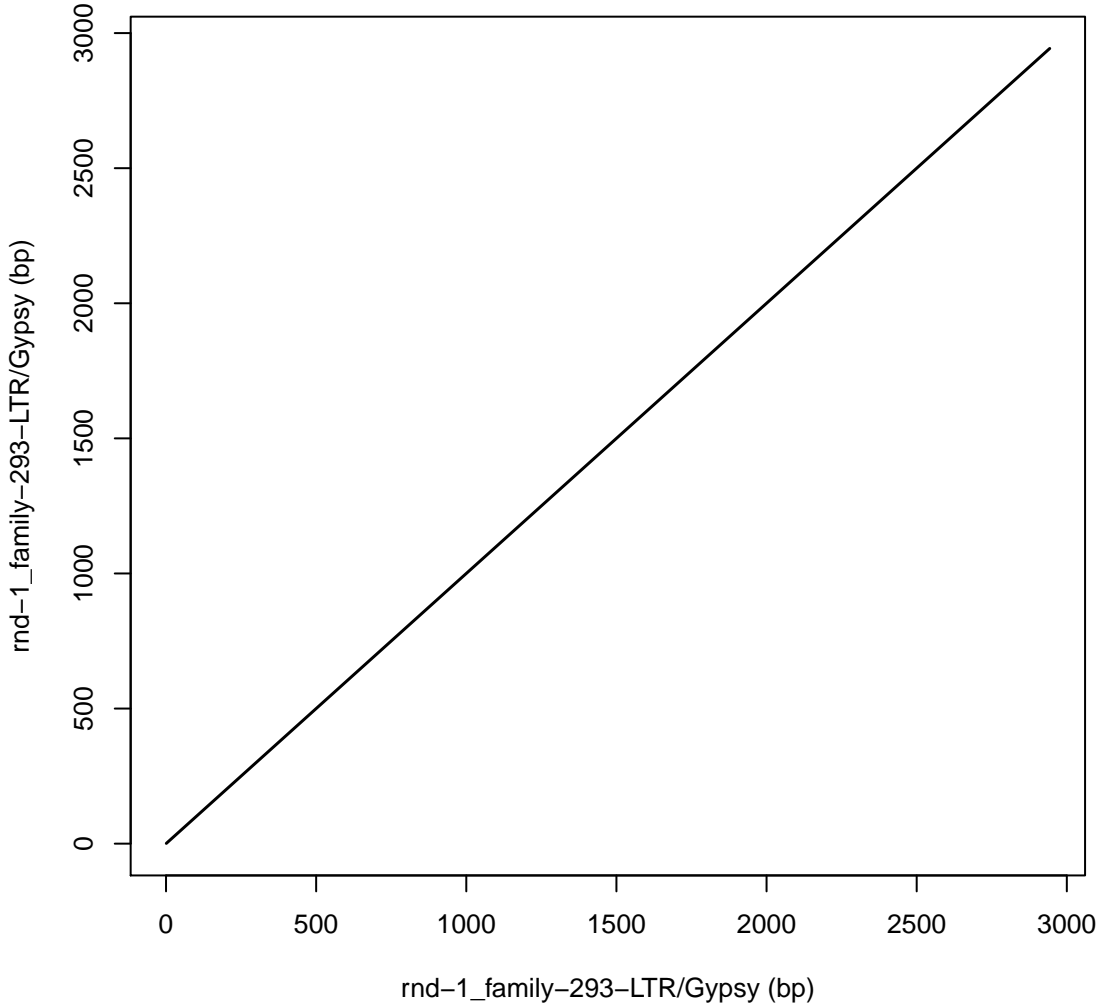
TE: rnd-1\_family-293-LTR/Gypsy  
consensus size: 2943bp; fragments: 377; full length: 20 ( $\geq 2648.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

