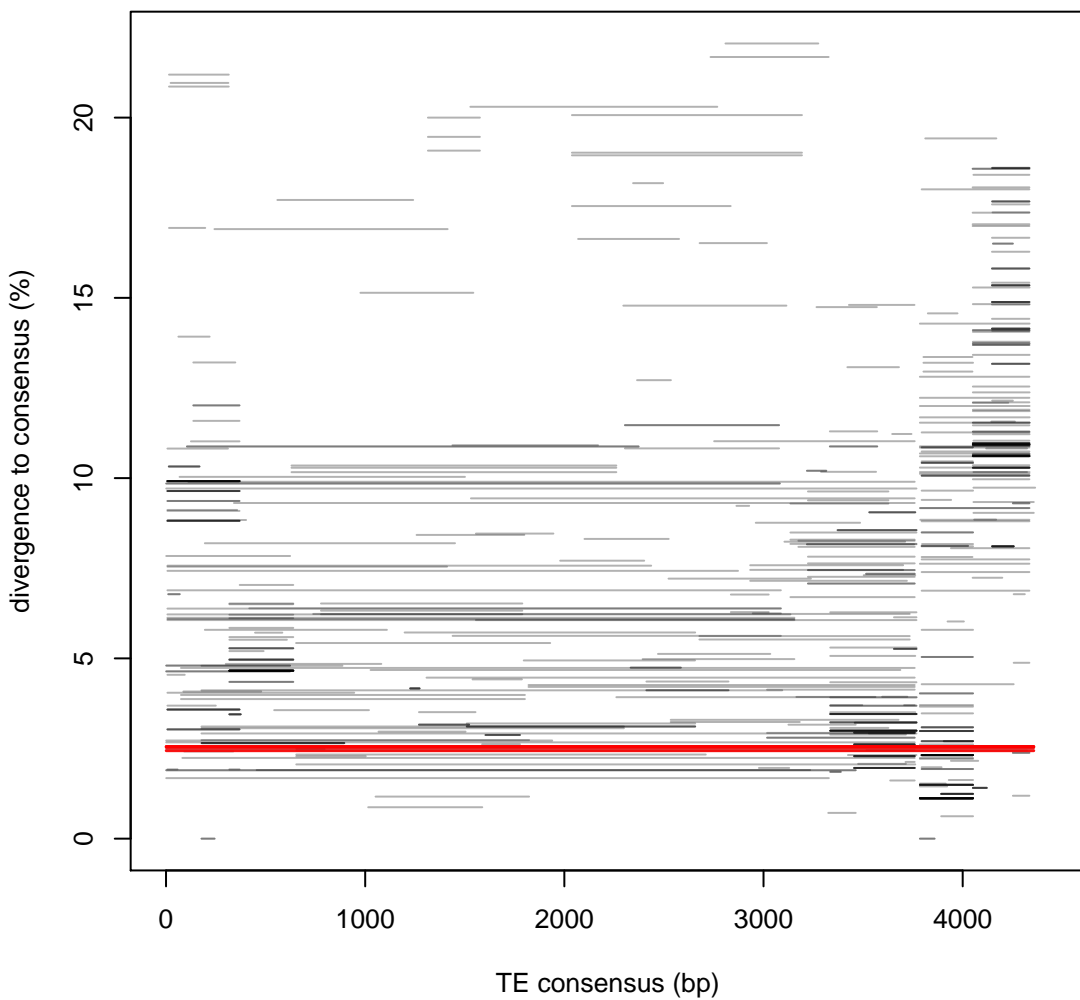
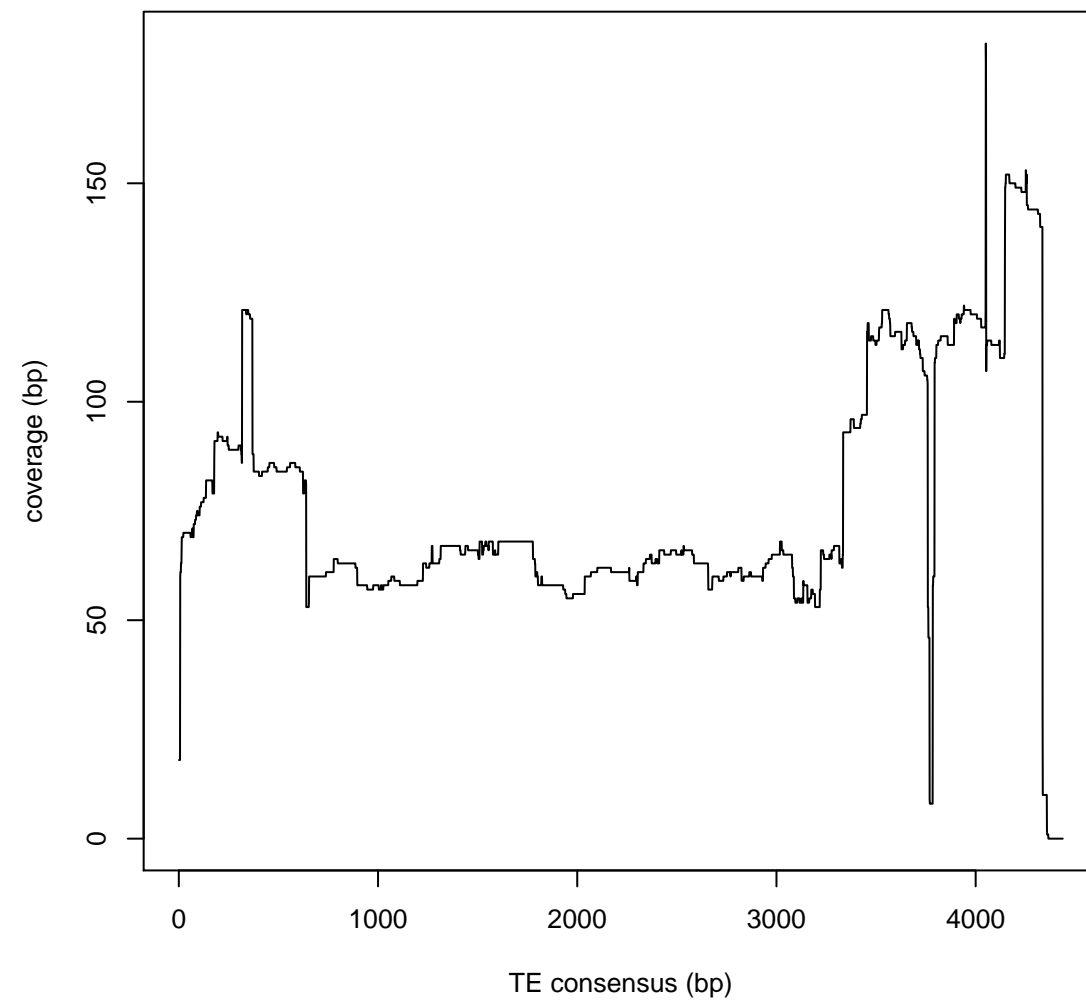


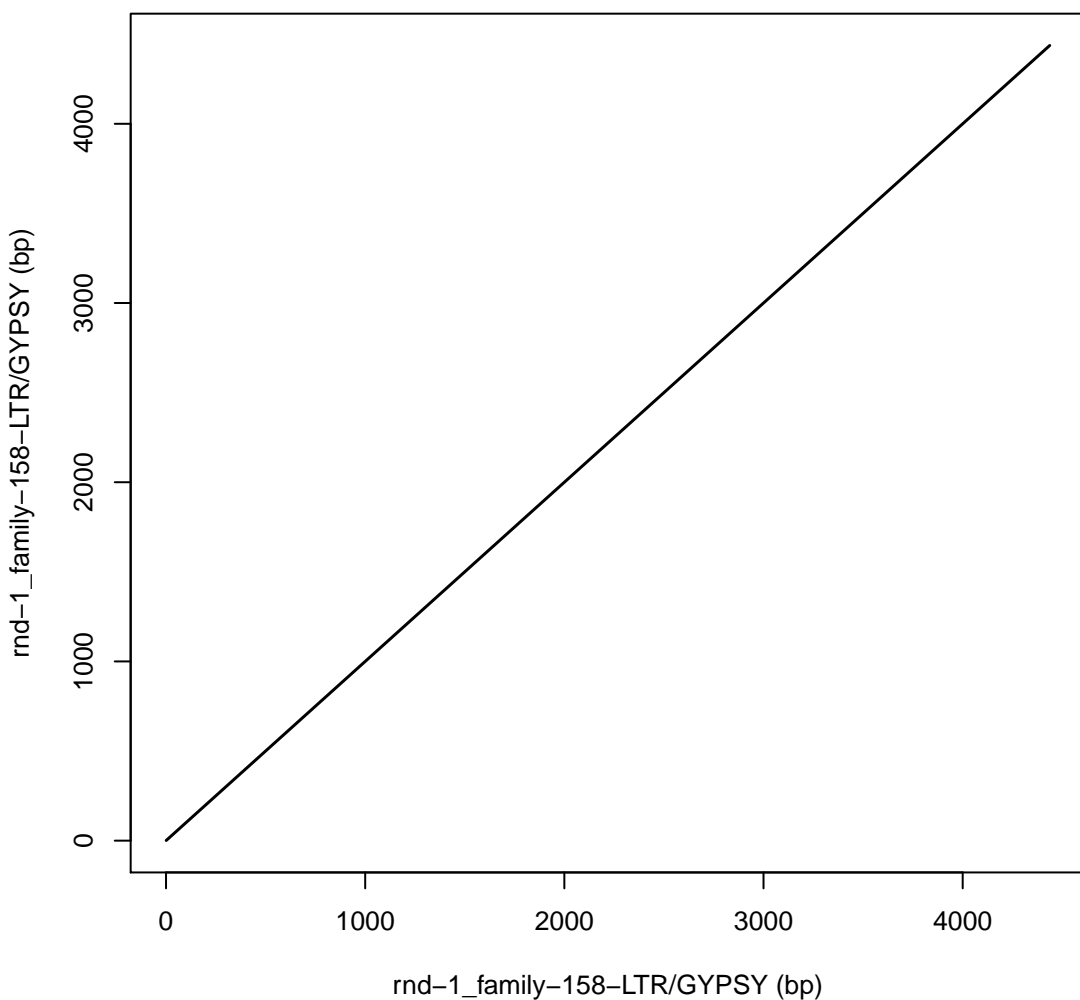
TE: rnd-1\_family-158-LTR/GYPSY  
consensus size: 4437bp; fragments: 587; full length: 4 ( $\geq 3993.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

