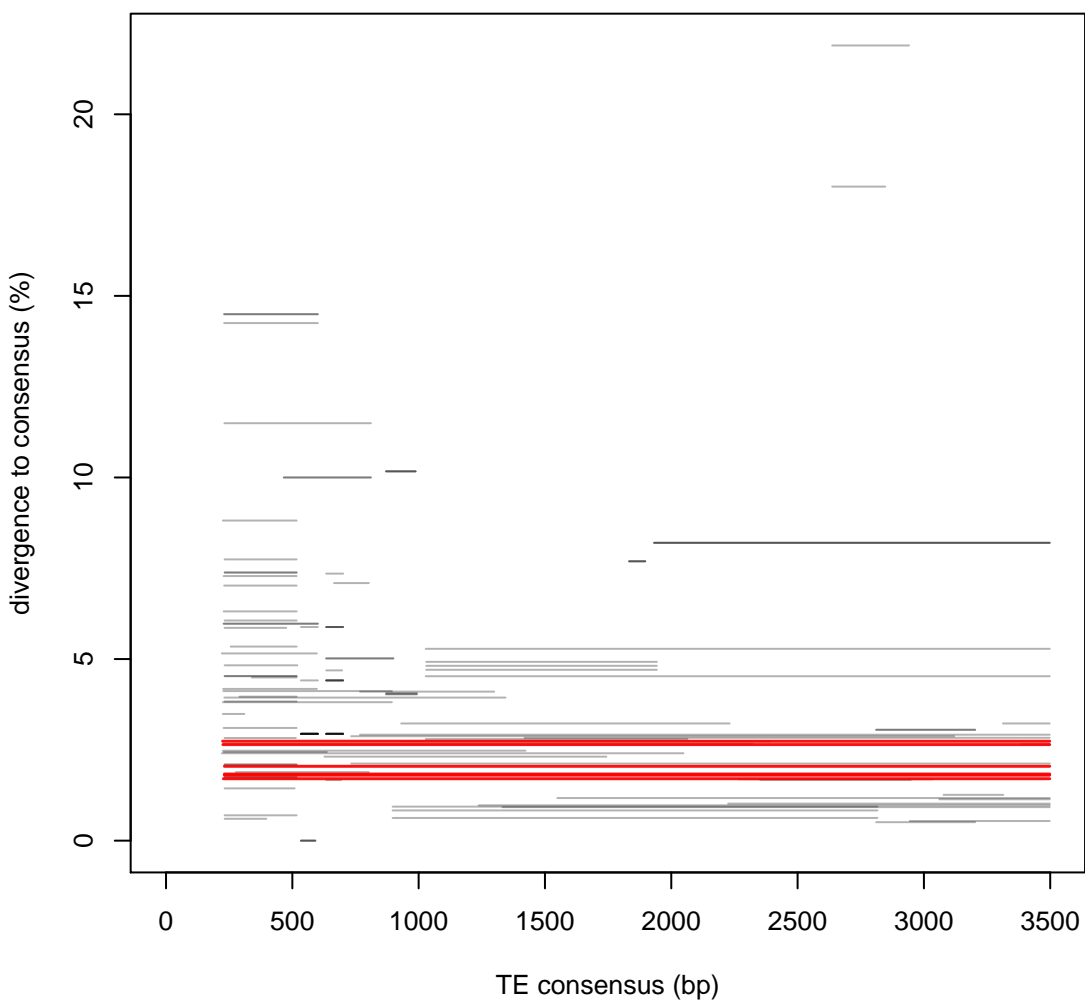
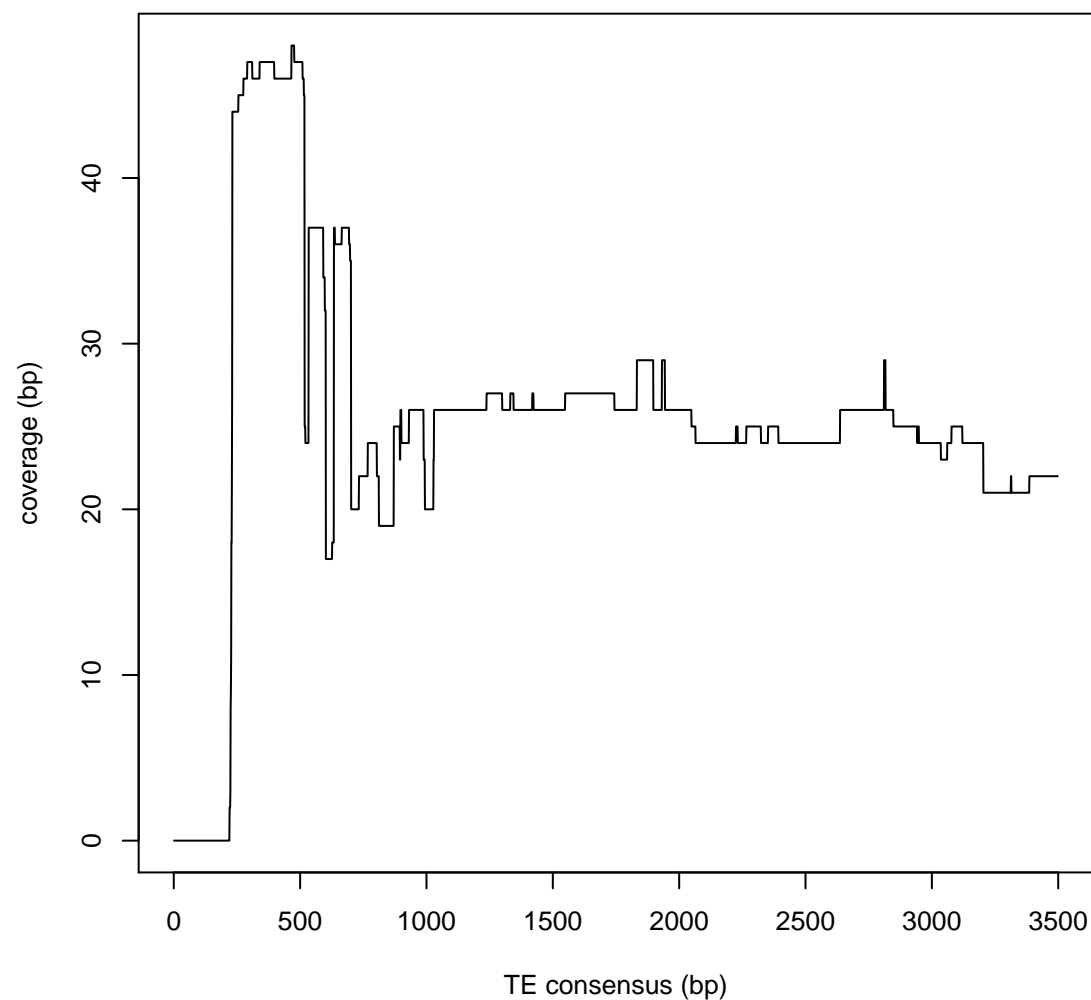


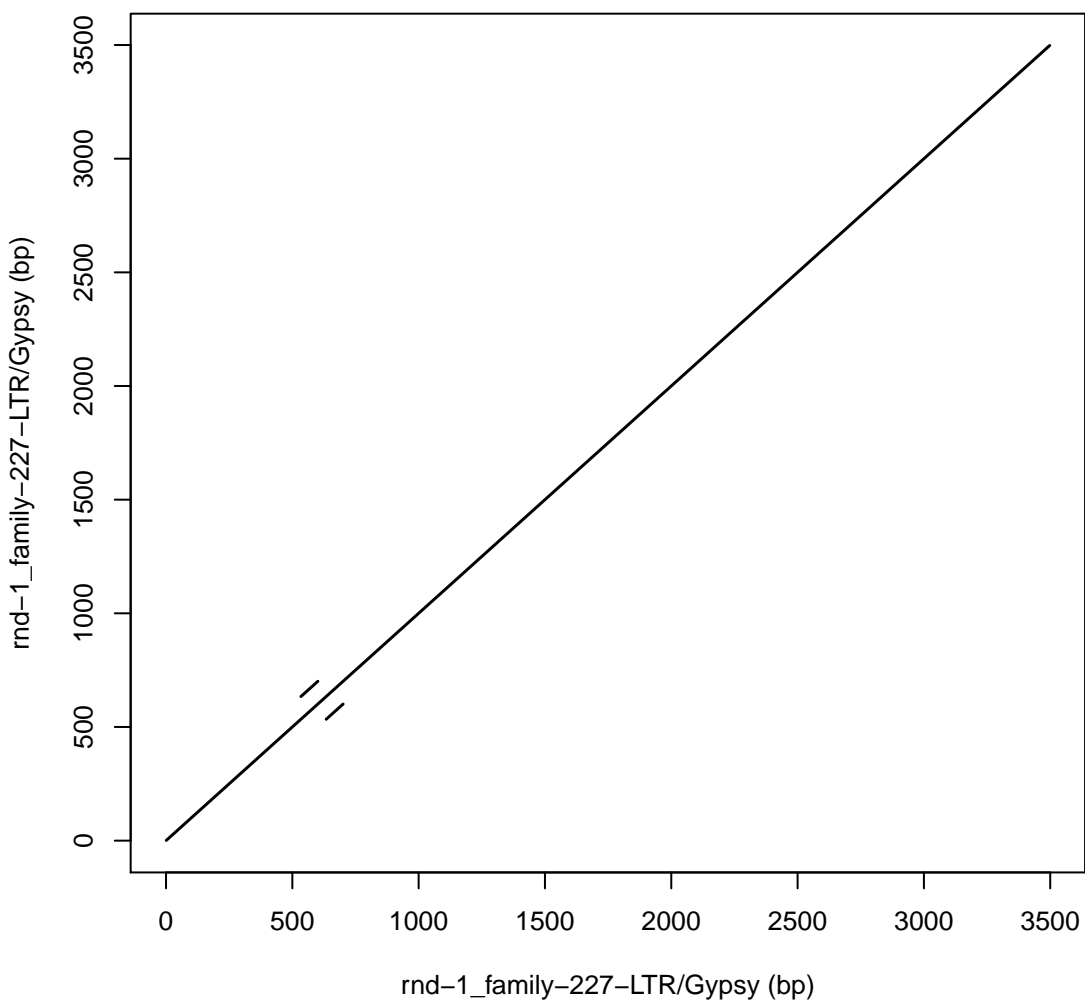
TE: rnd-1\_family-227-LTR/Gypsy  
consensus size: 3498bp; fragments: 128; full length: 6 ( $\geq 3148.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

