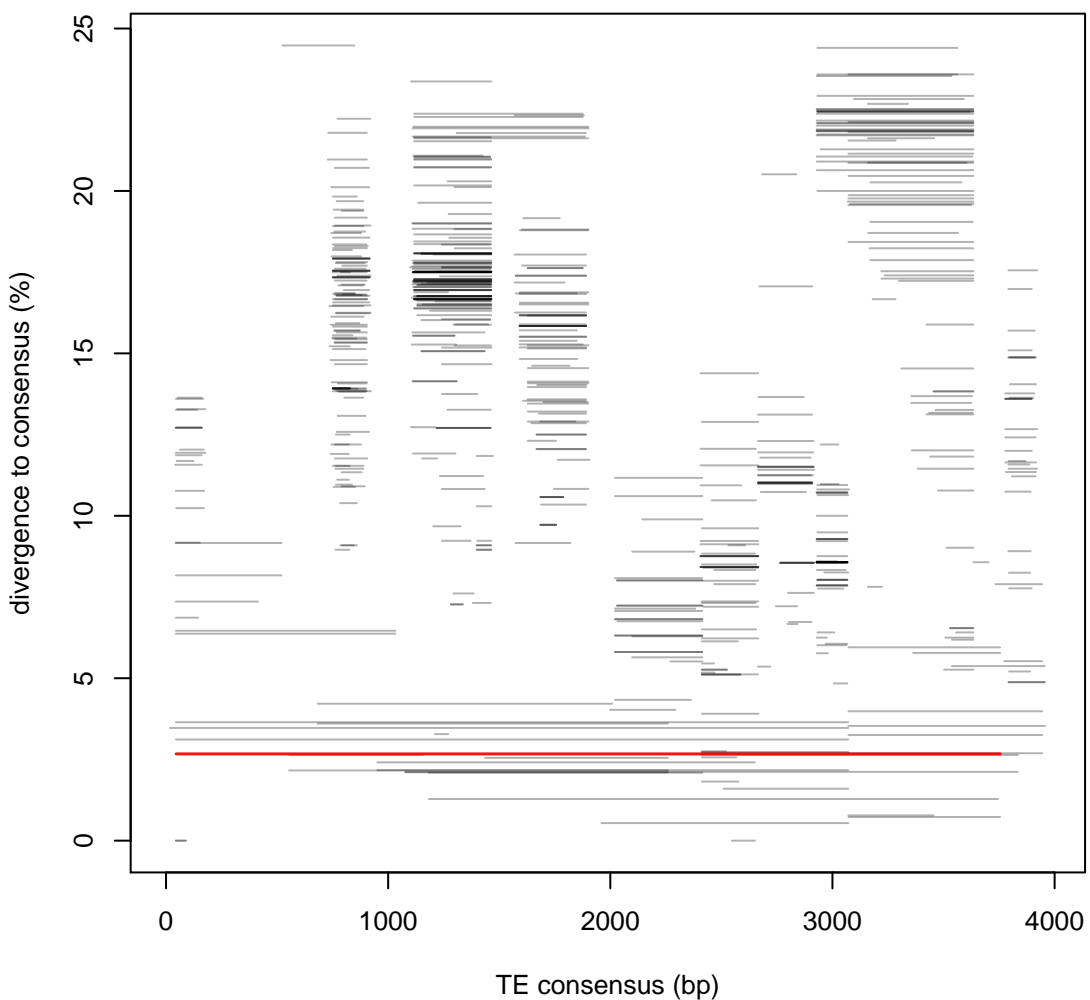
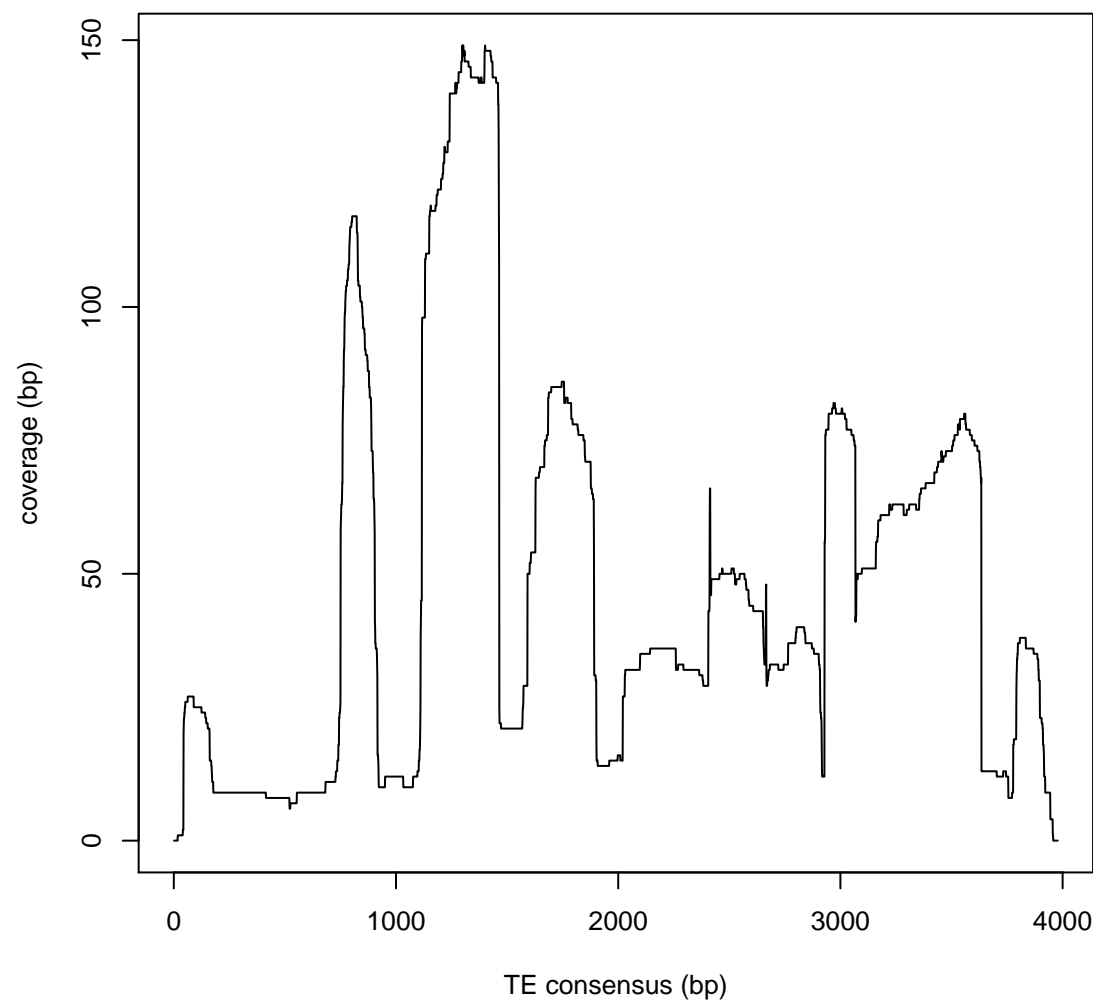


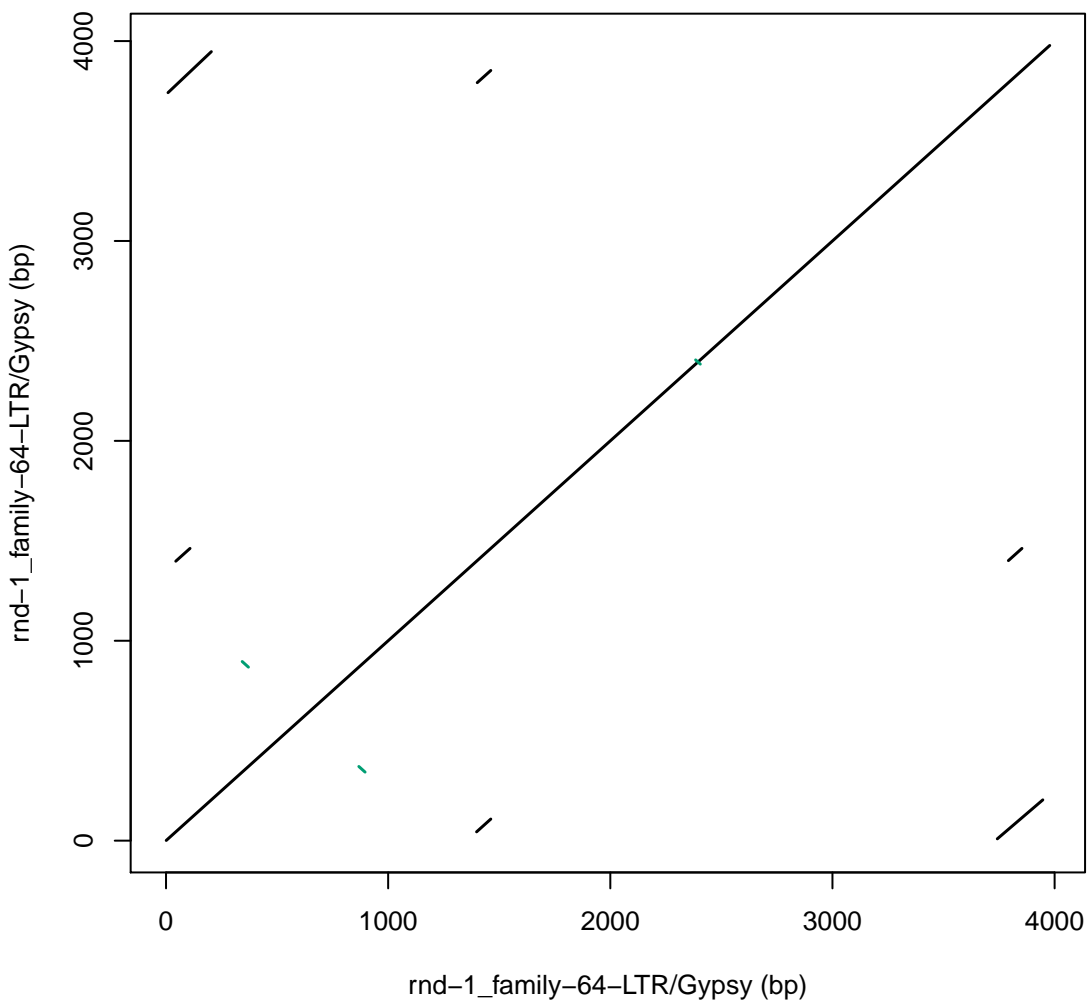
TE: rnd-1\_family-64-LTR/Gypsy  
consensus size: 3978bp; fragments: 621; full length: 1 ( $\geq 3580.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

