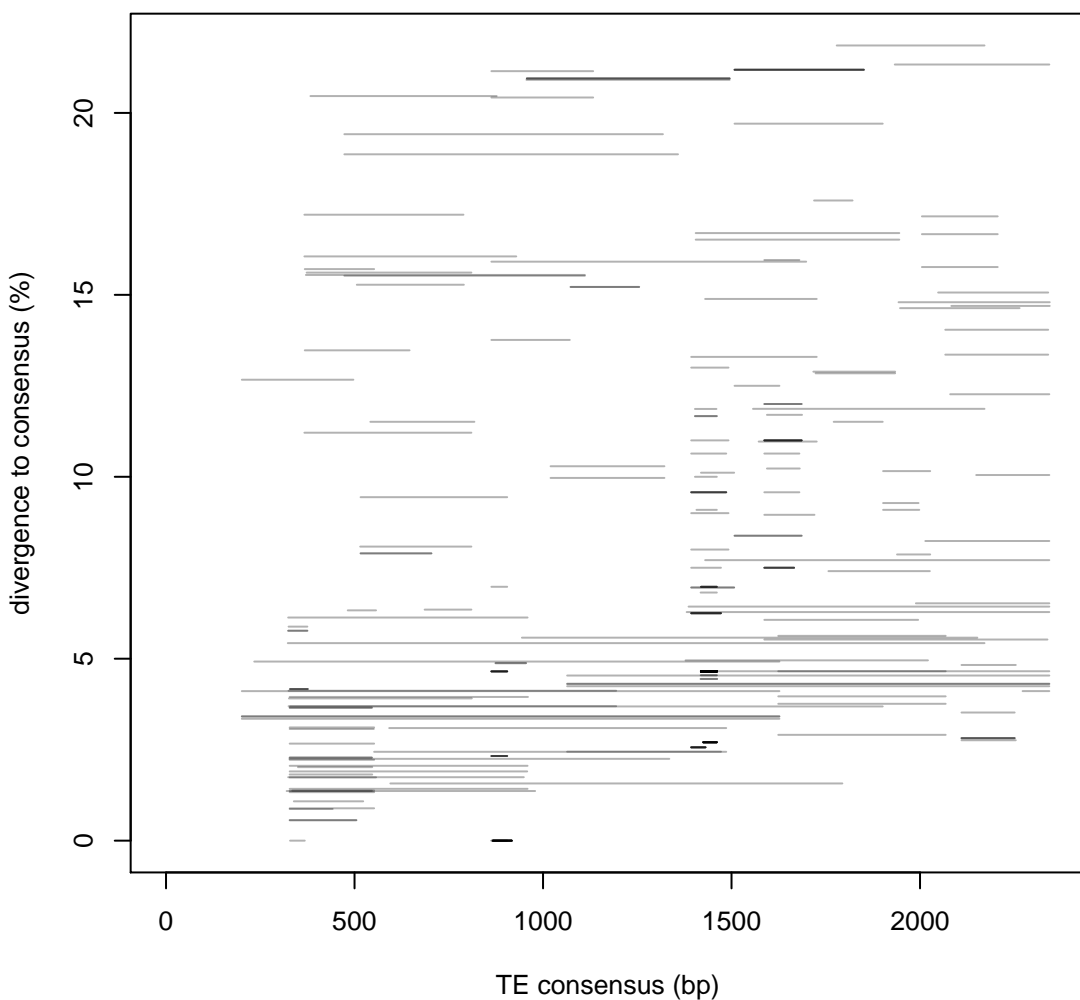
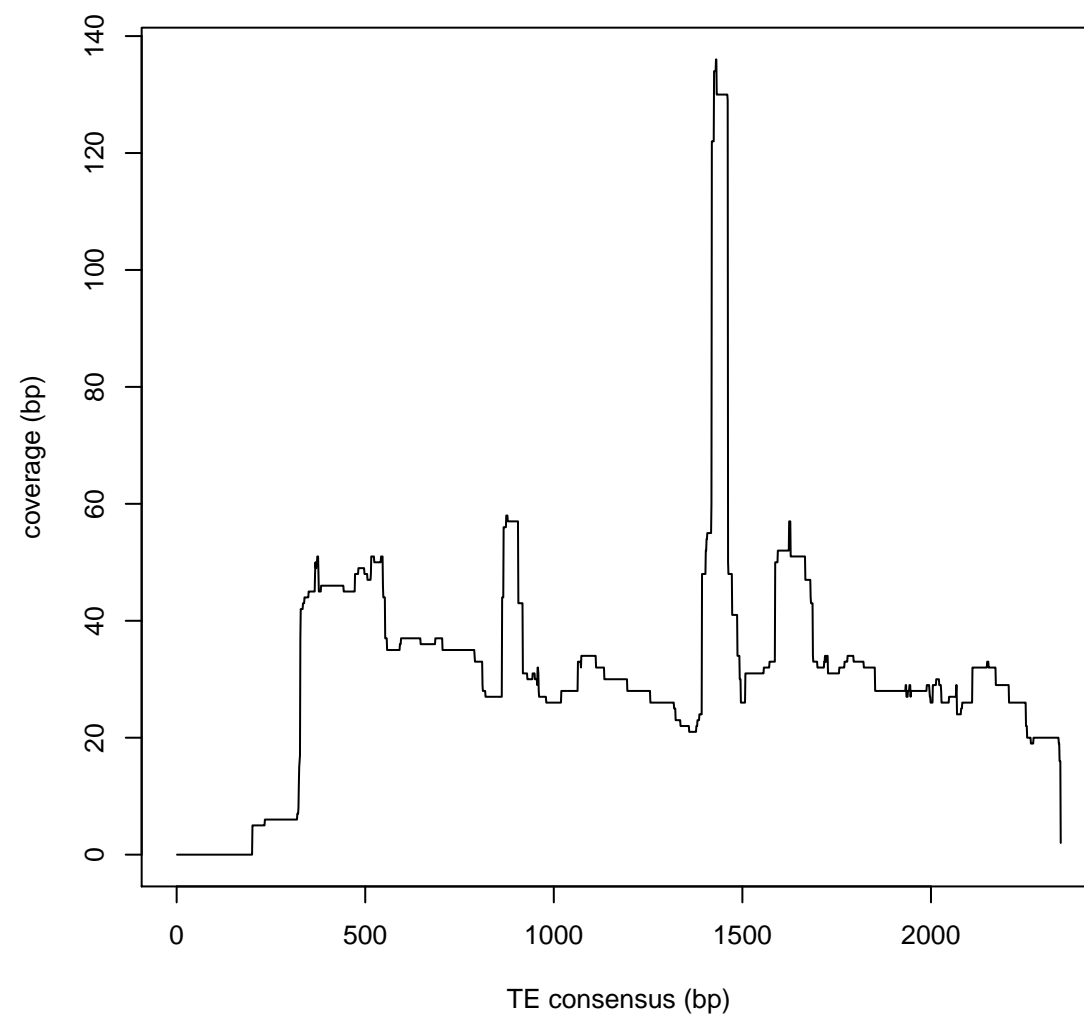


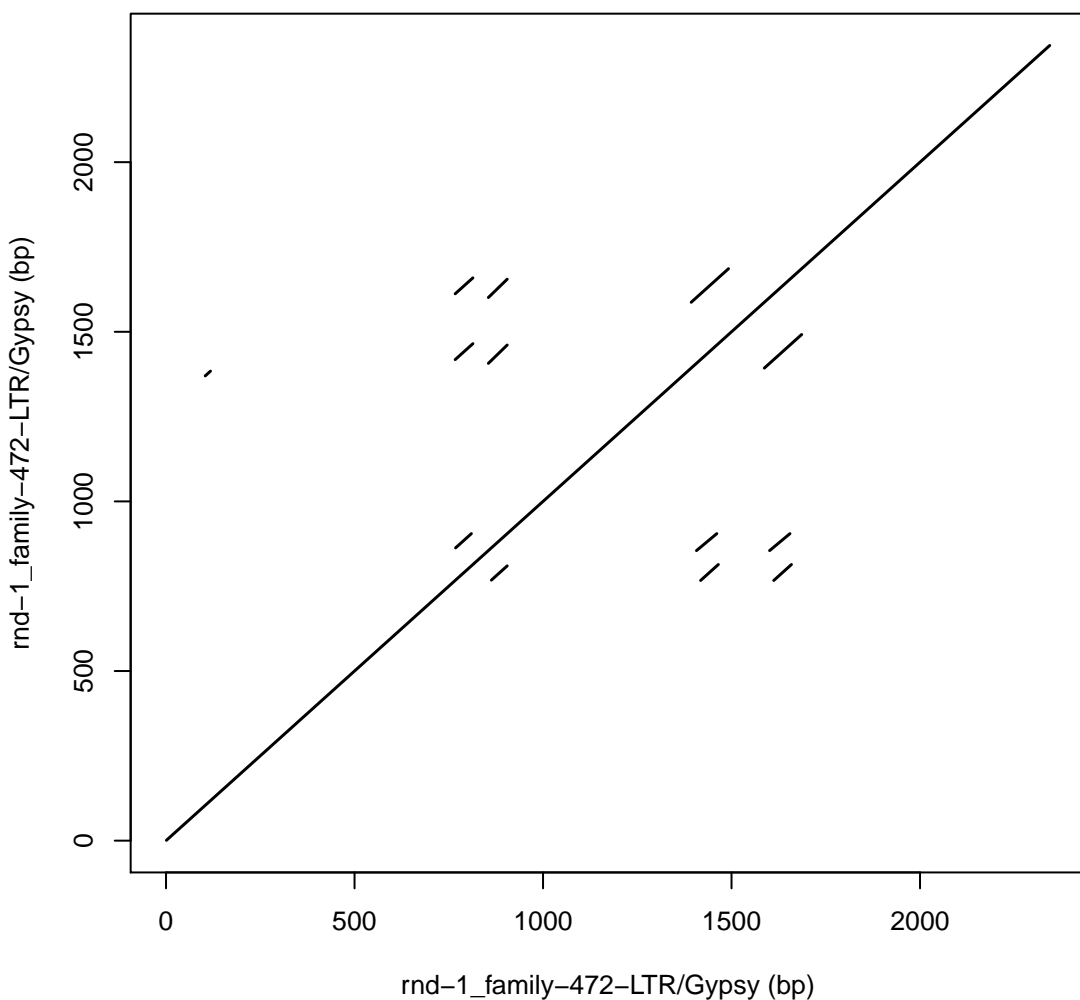
TE: rnd-1\_family-472-LTR/Gypsy  
consensus size: 2344bp; fragments: 292; full length: 0 ( $\geq 2109.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

