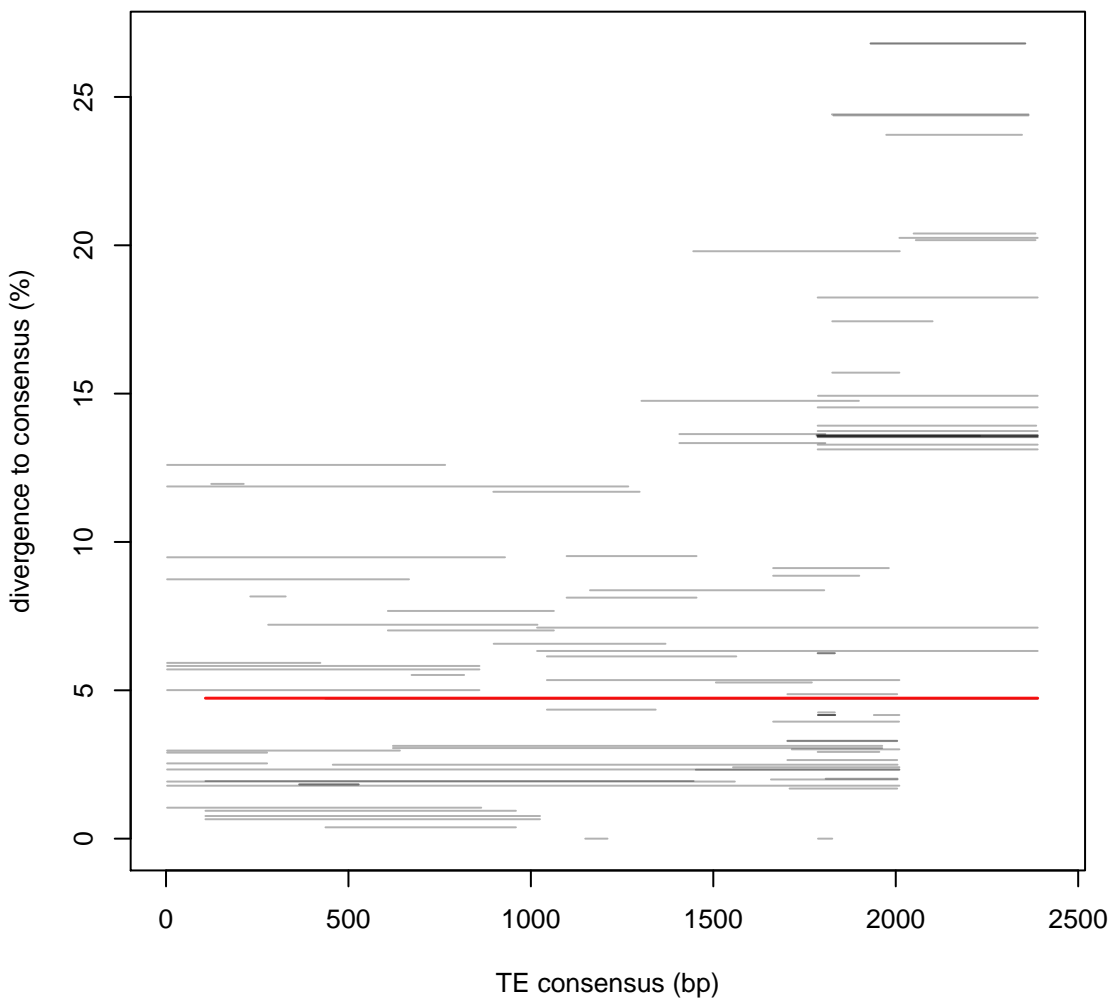
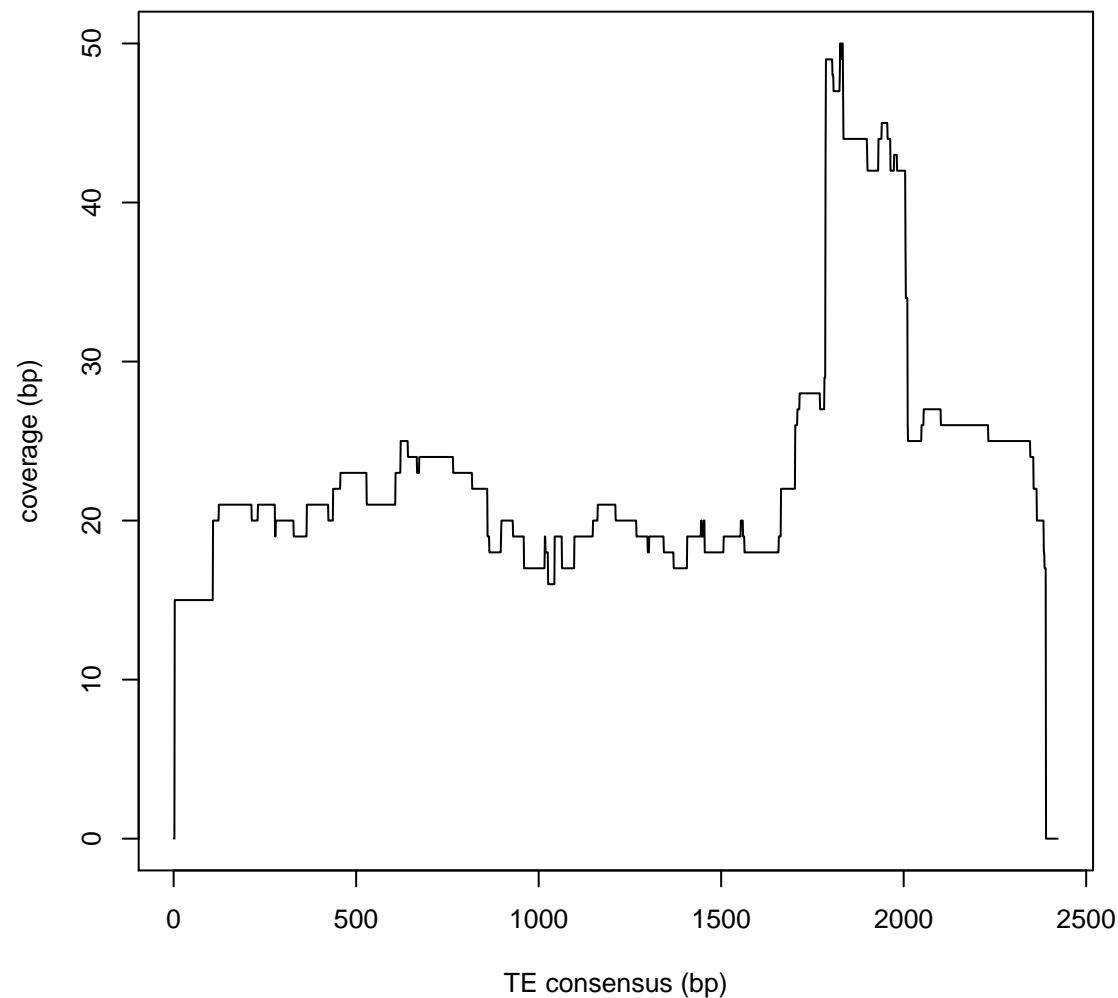


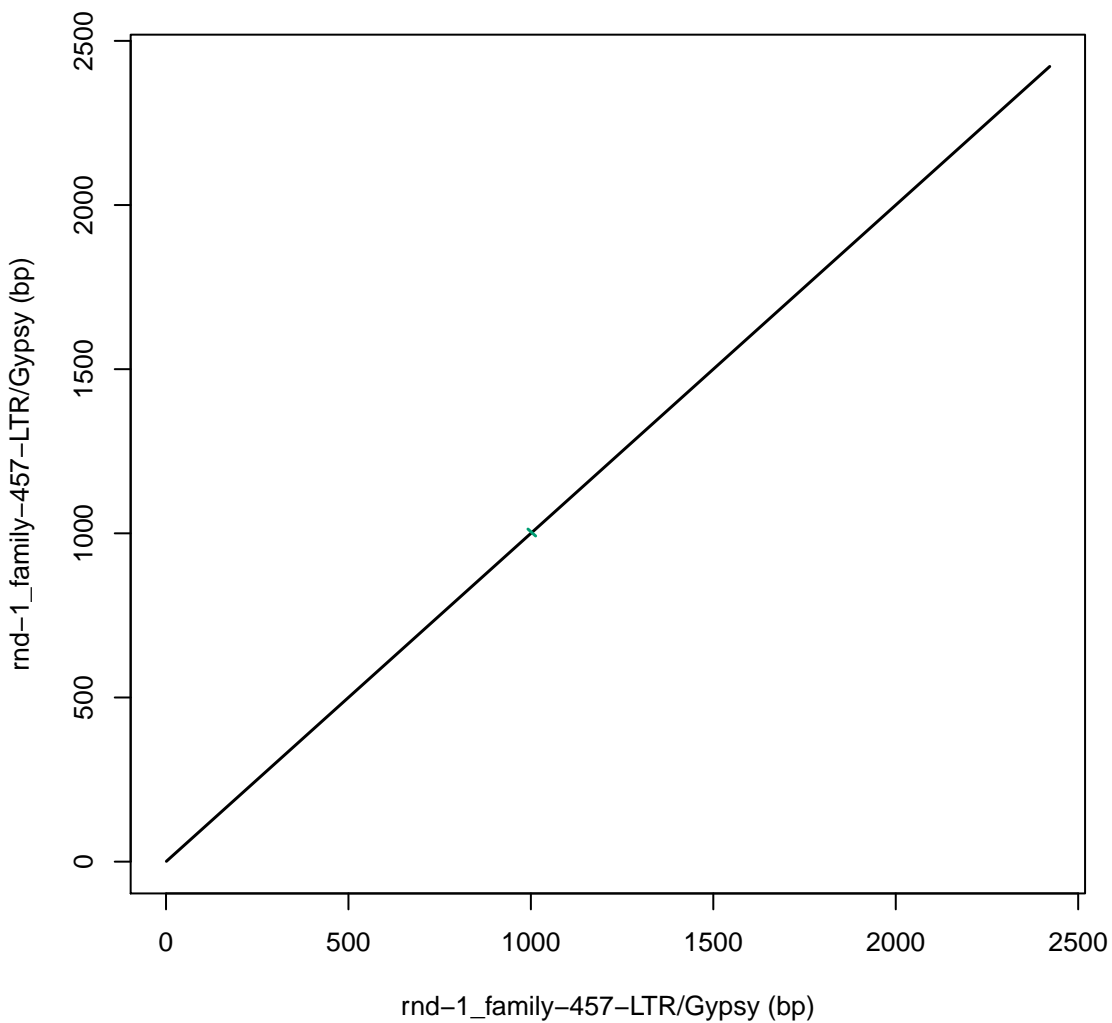
TE: rnd-1\_family-457-LTR/Gypsy  
consensus size: 2422bp; fragments: 95; full length: 1 ( $\geq 2179.8\text{bp}$ )



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

