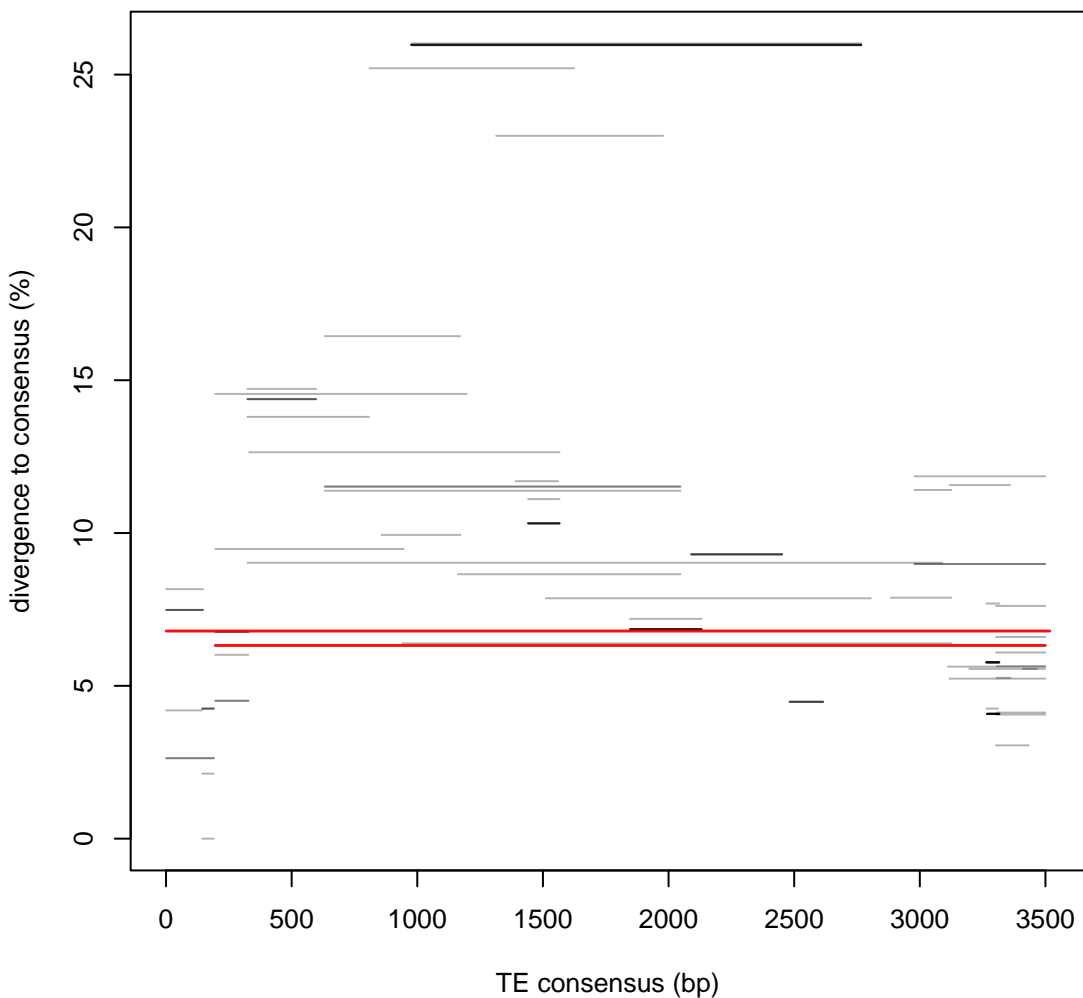
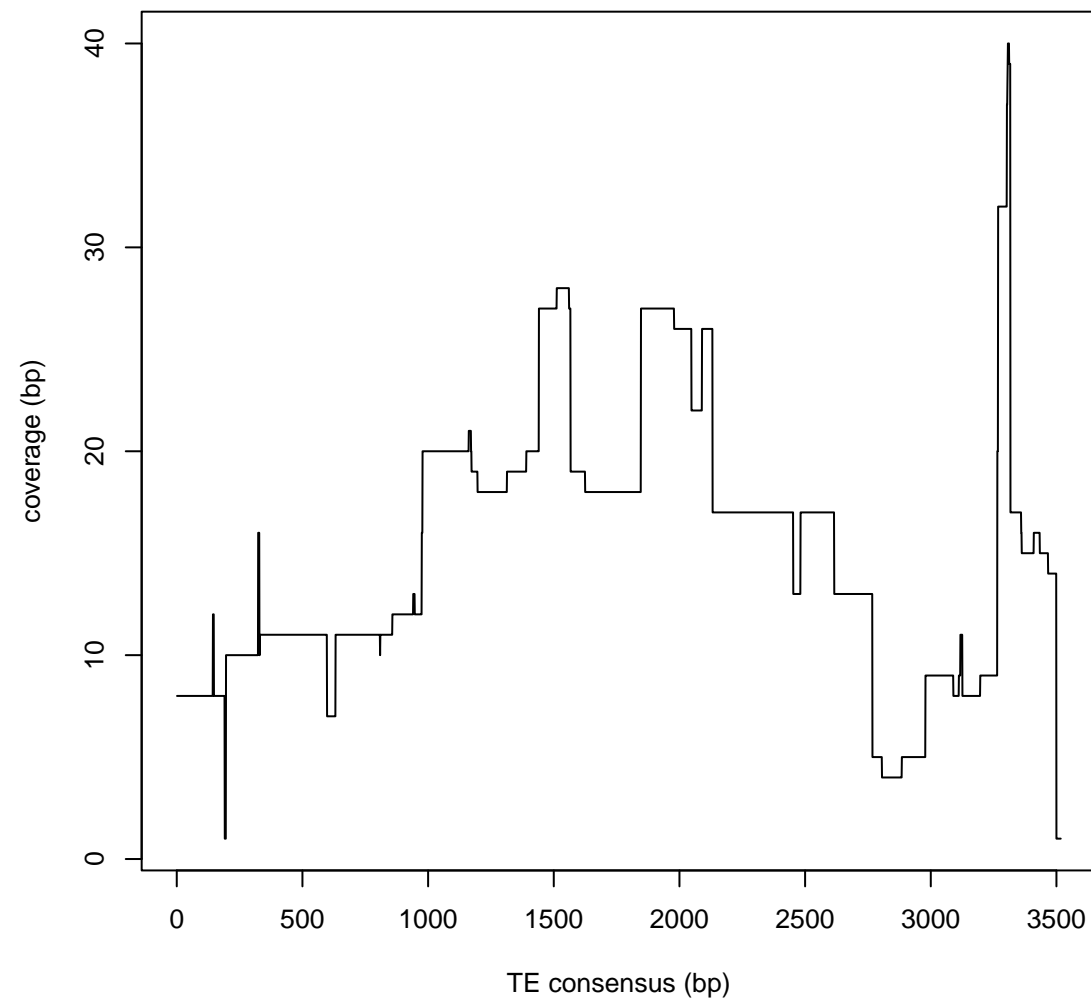


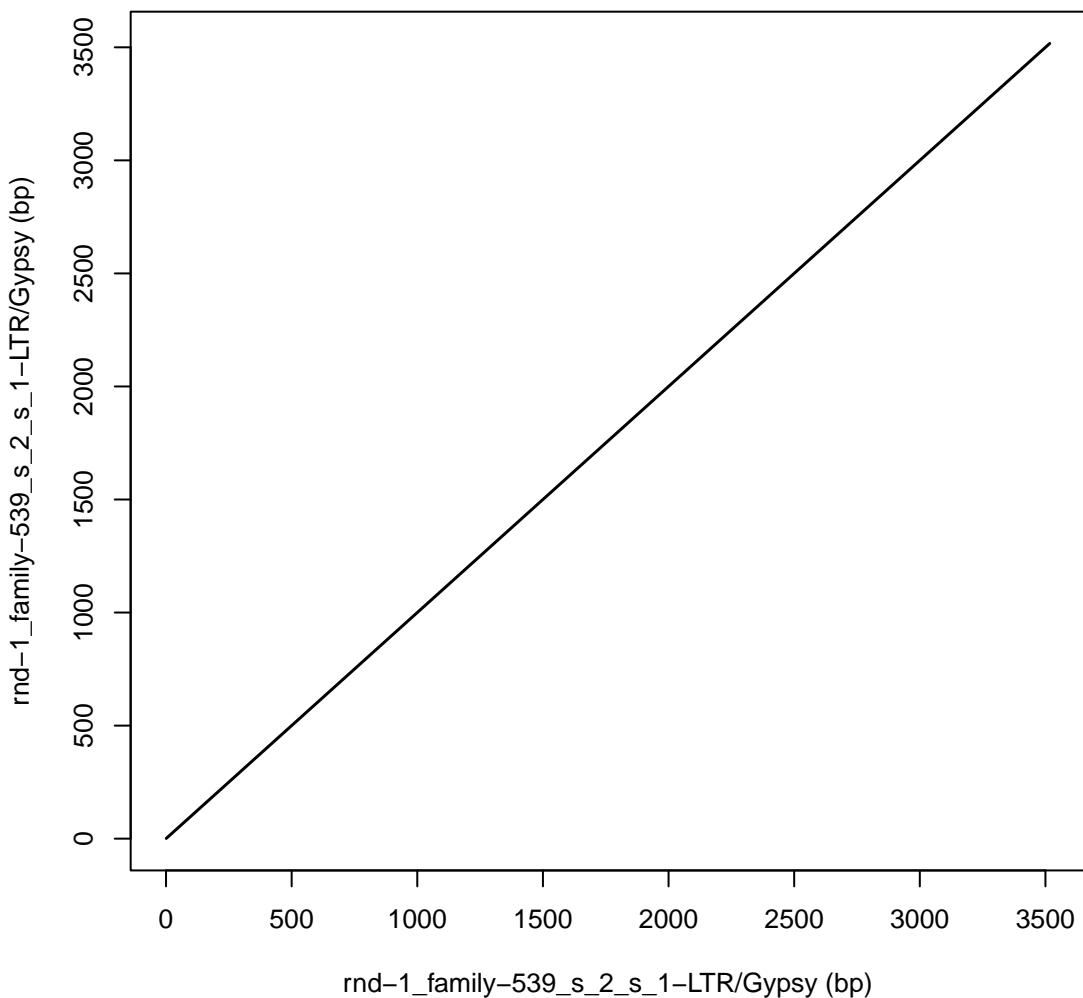
TE: rnd-1\_family-539\_s\_2\_s\_1-LTR/Gypsy  
consensus size: 3517bp; fragments: 113; full length: 2 ( $\geq 3165.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

