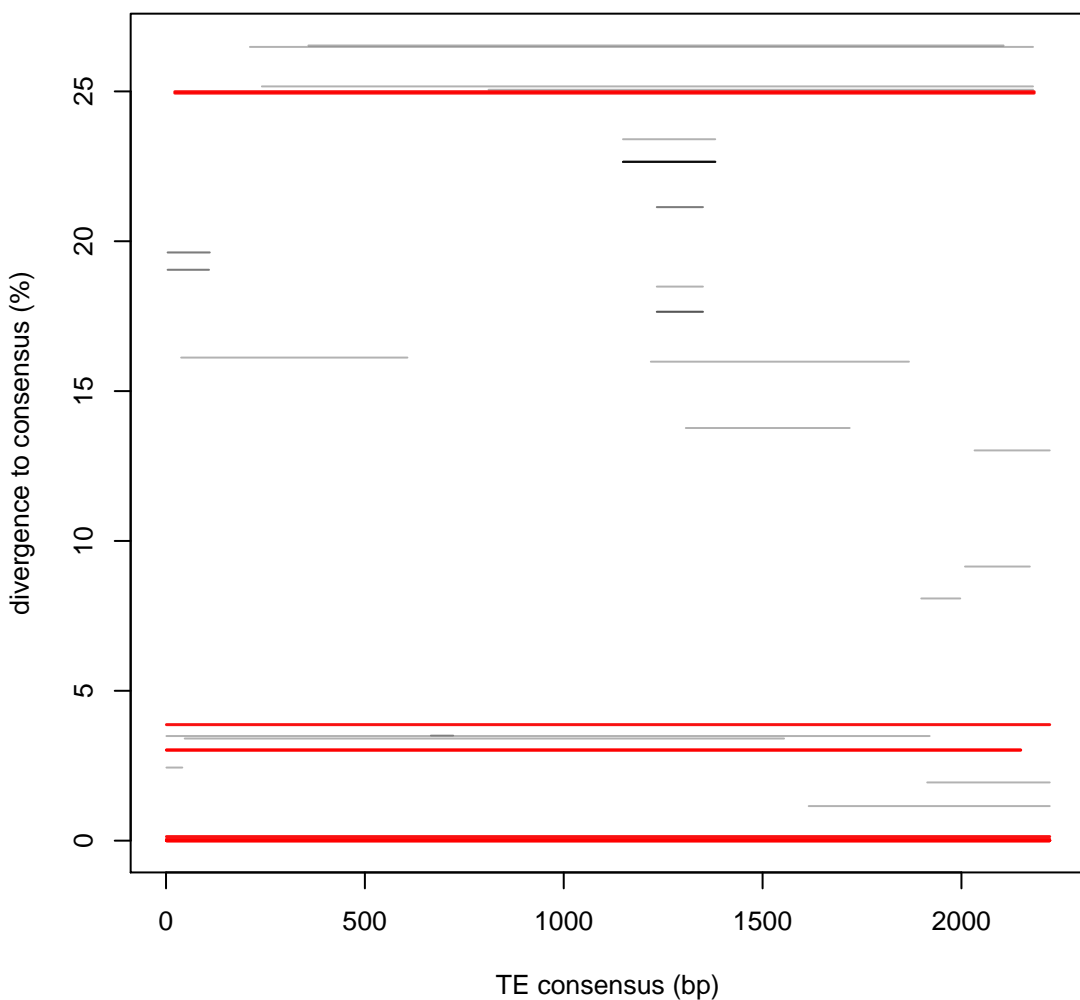
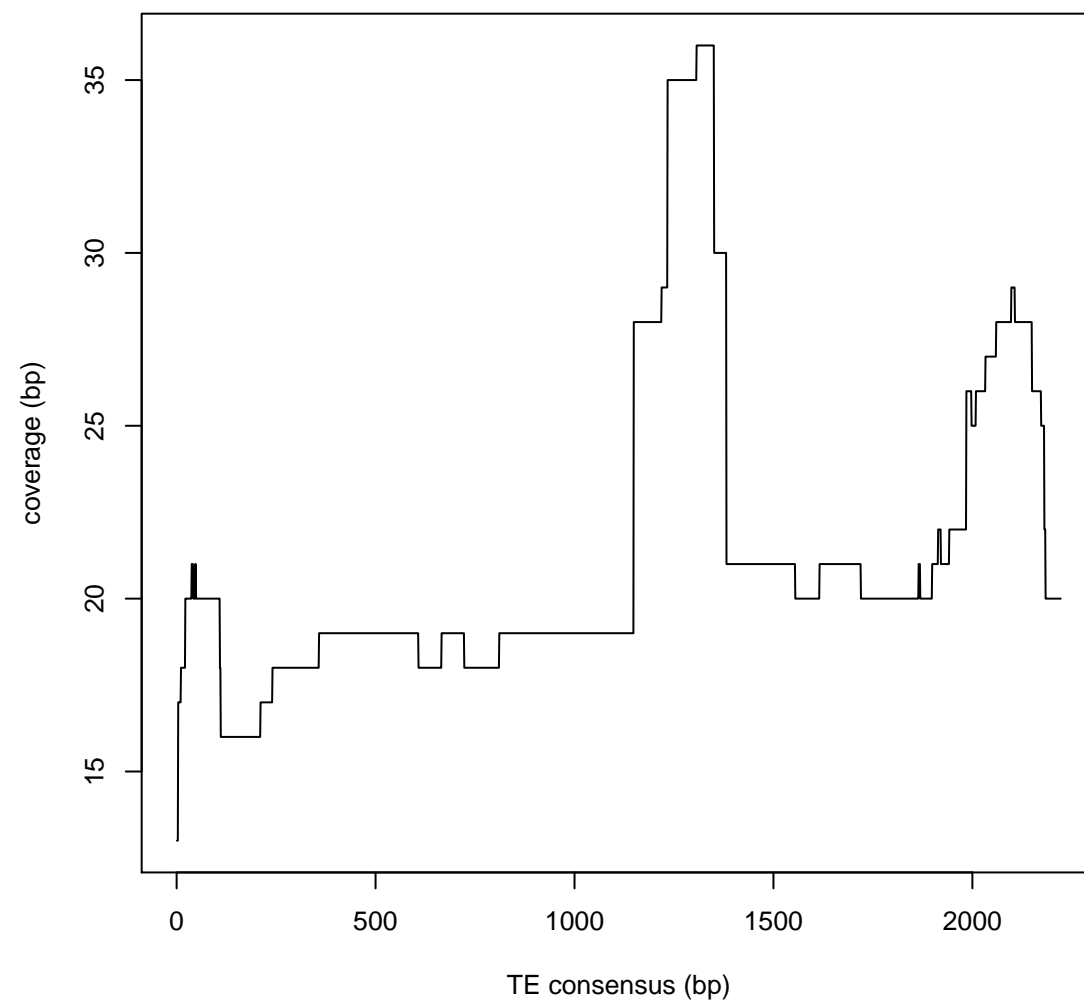


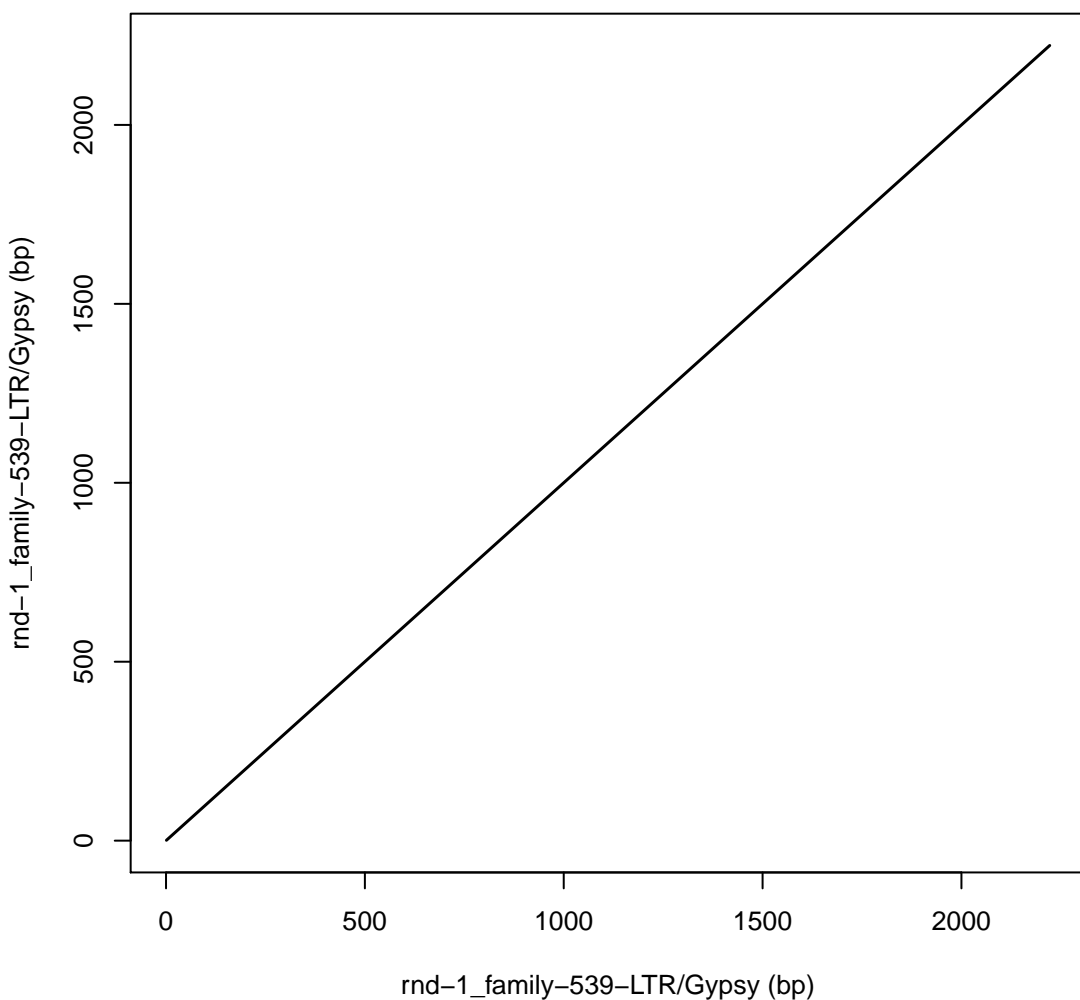
TE: rnd-1\_family-539-LTR/Gypsy  
consensus size: 2222bp; fragments: 57; full length: 13 ( $\geq 1999.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

