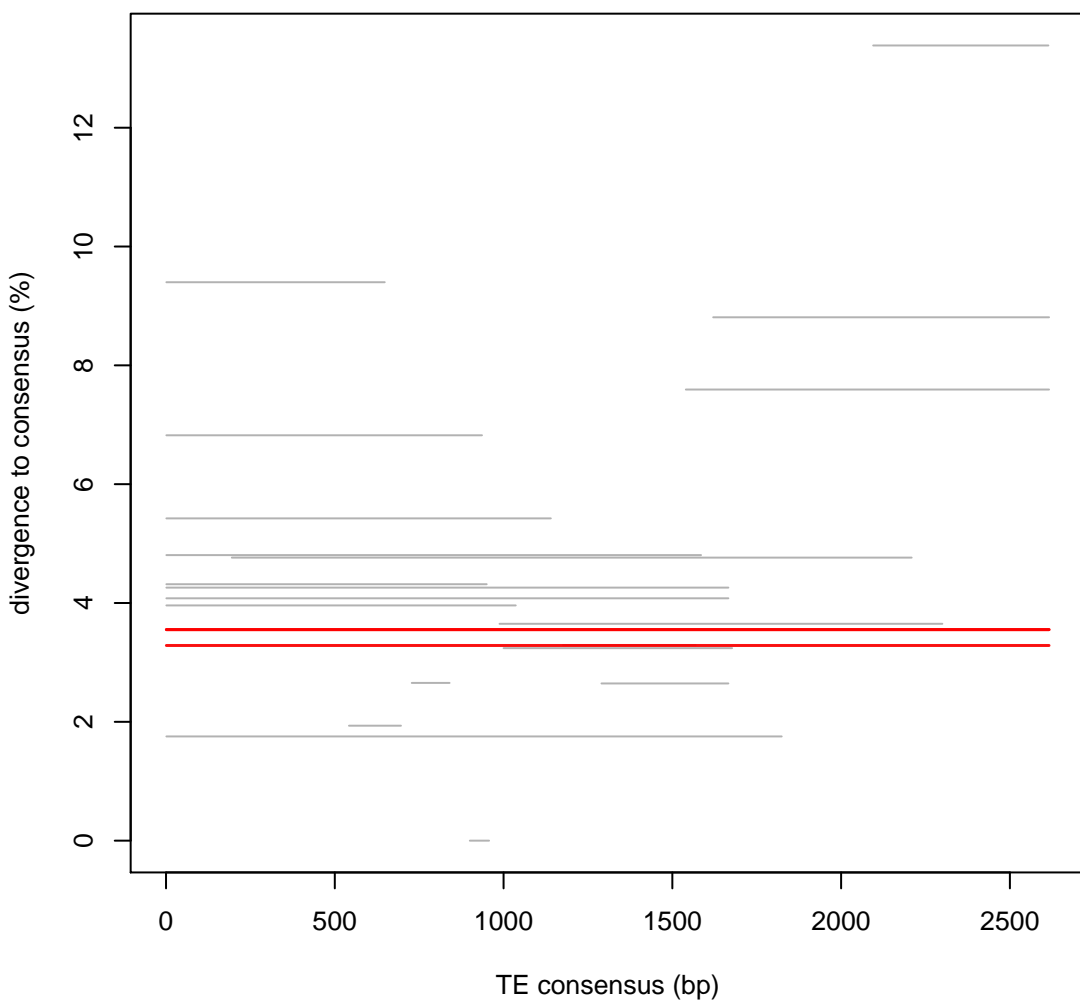
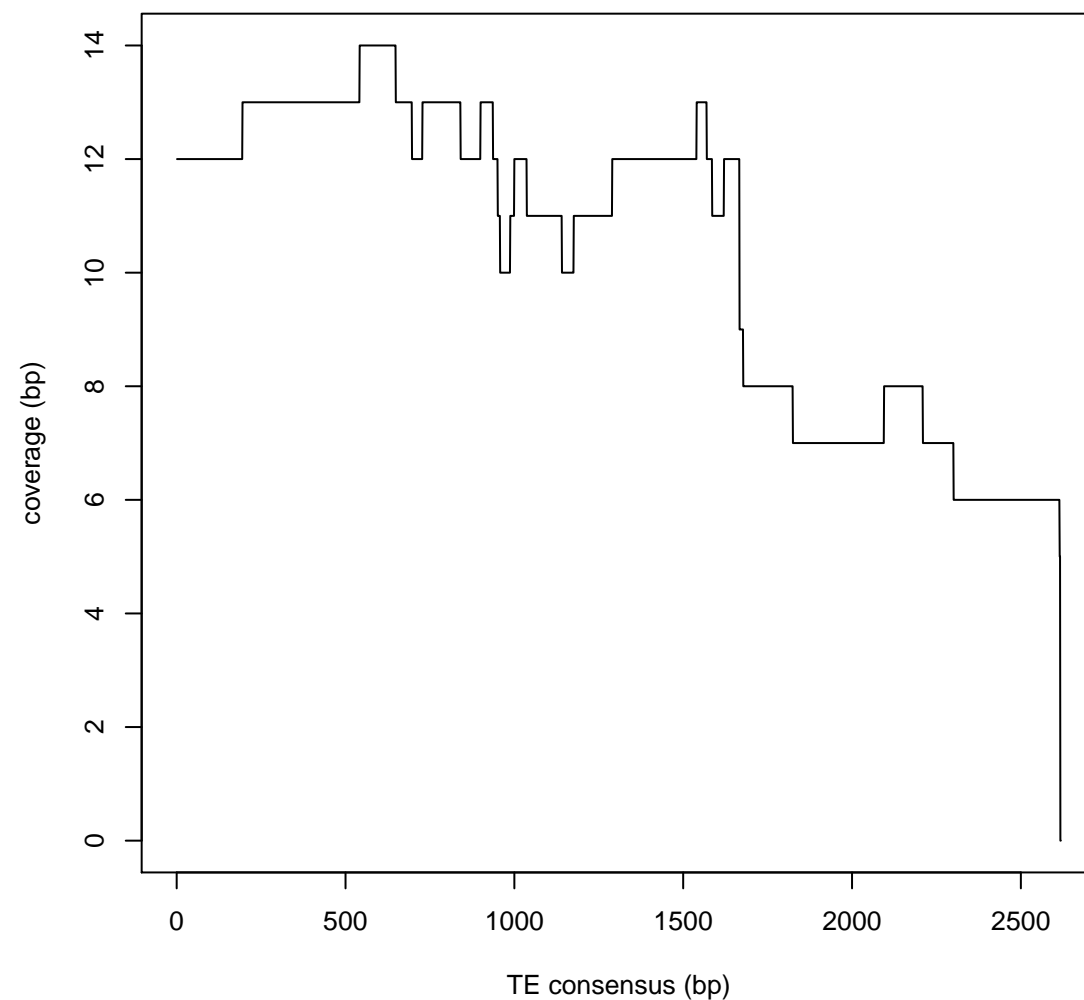


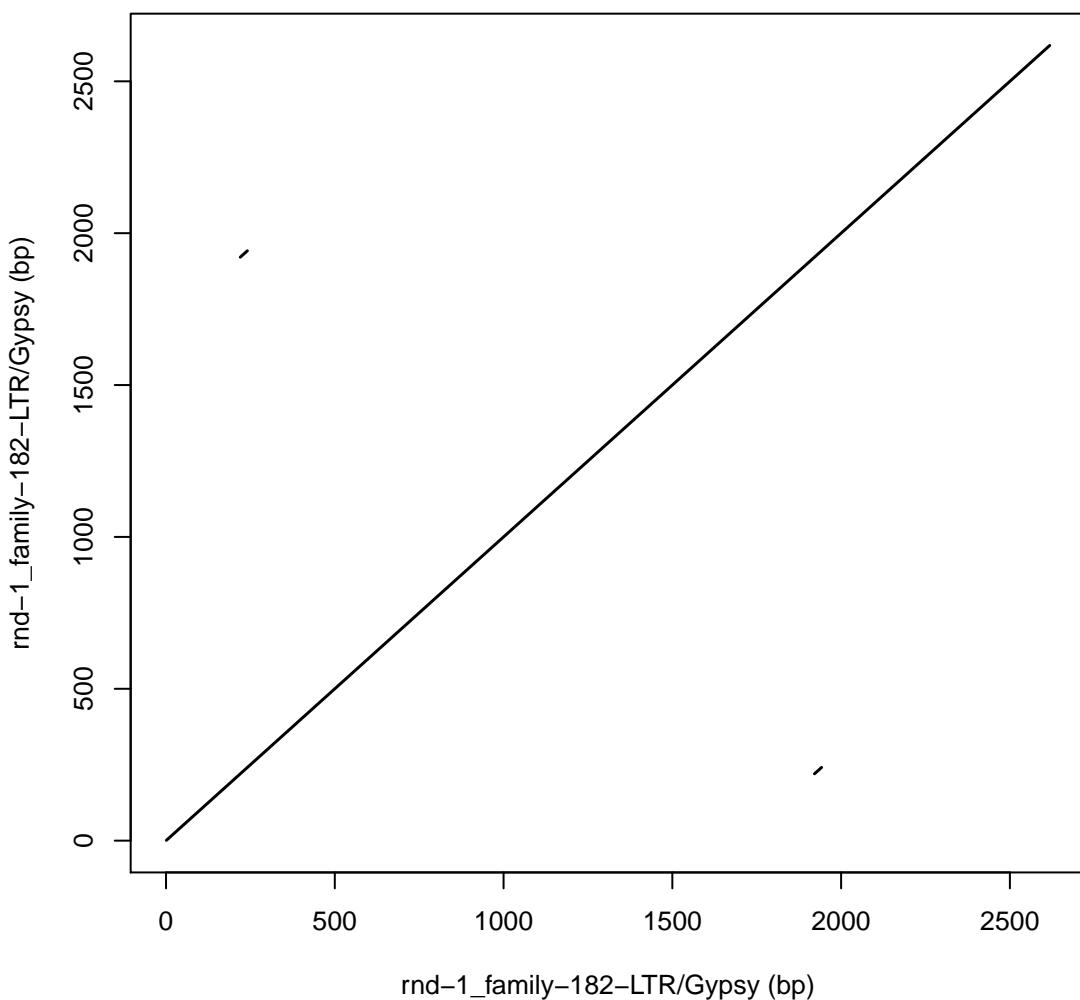
TE: rnd-1_family-182-LTR/Gypsy
consensus size: 2618bp; fragments: 23; full length: 3 ($\geq 2356.2\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

