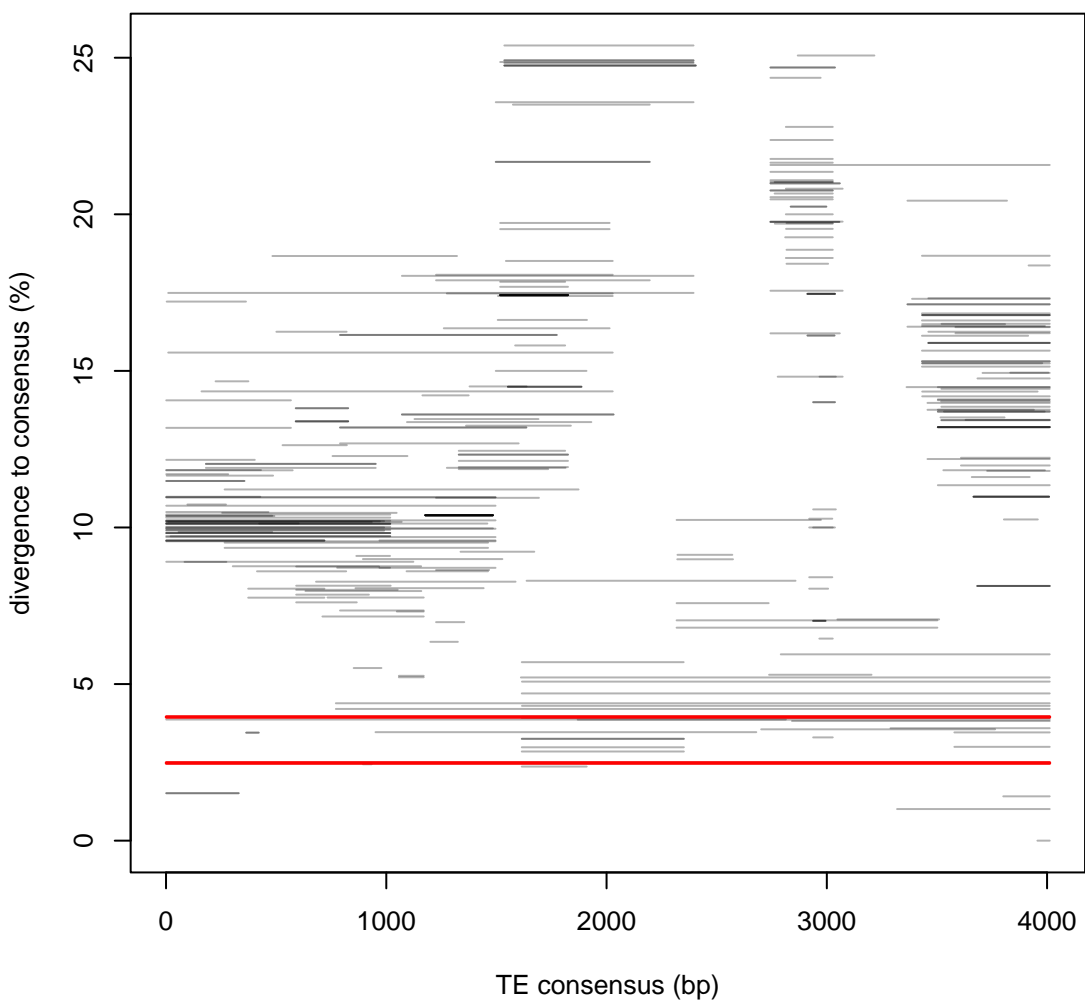
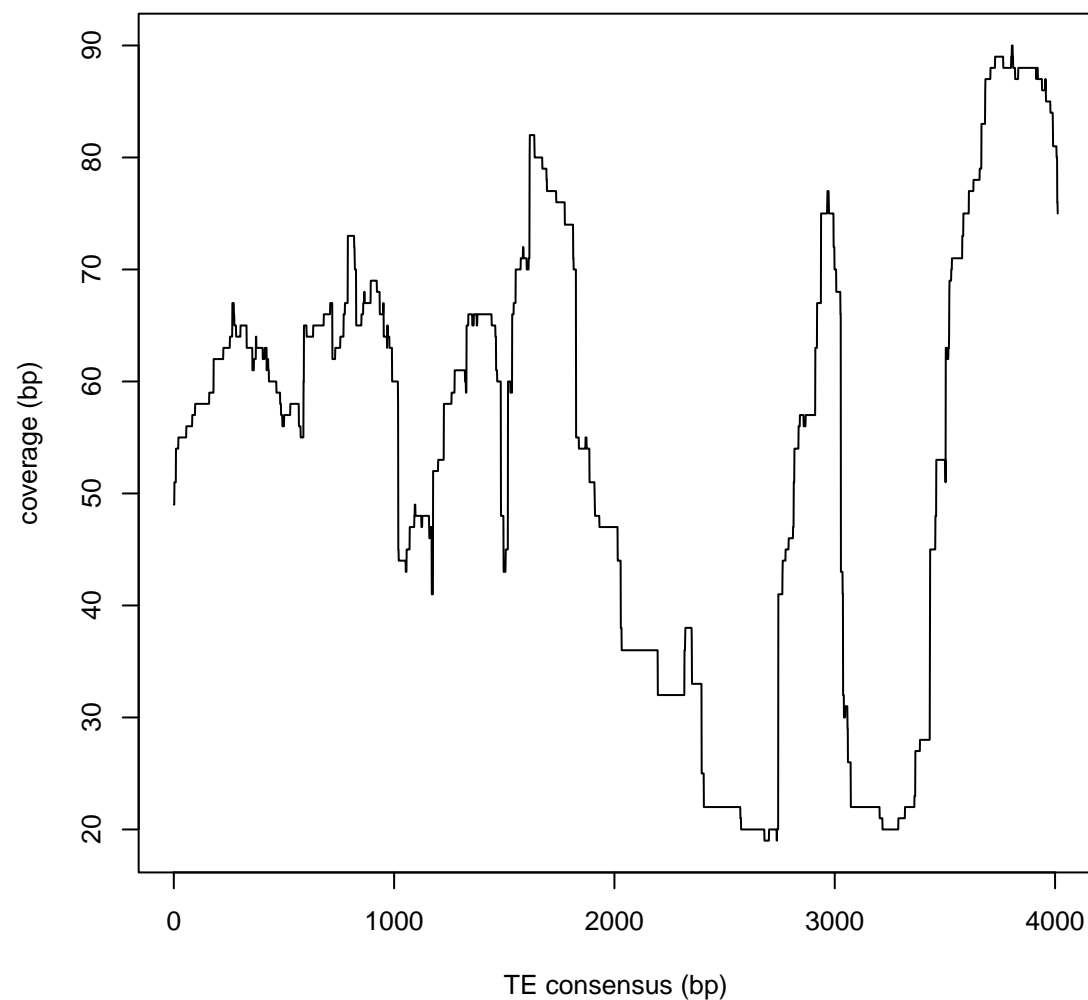


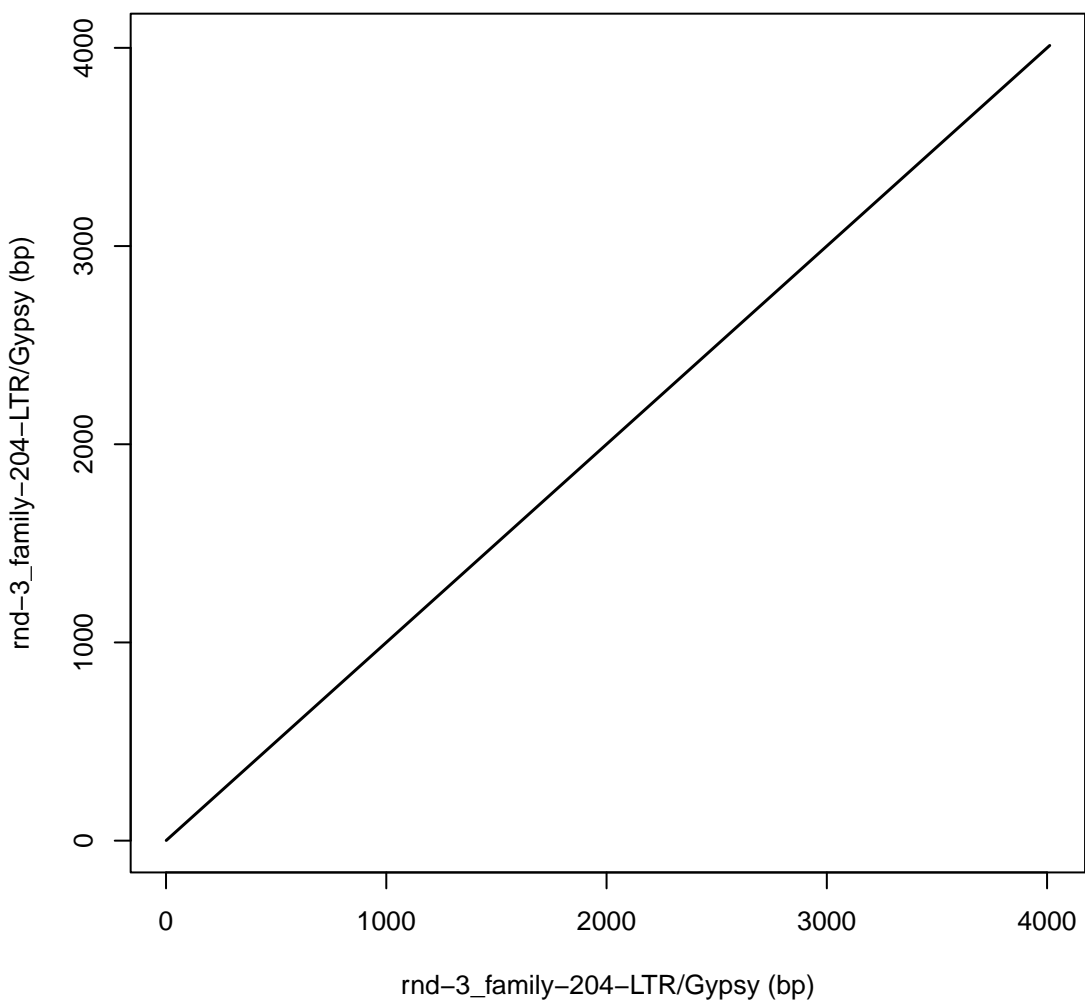
TE: rnd-3\_family-204-LTR/Gypsy  
consensus size: 4012bp; fragments: 341; full length: 4 ( $\geq 3610.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

