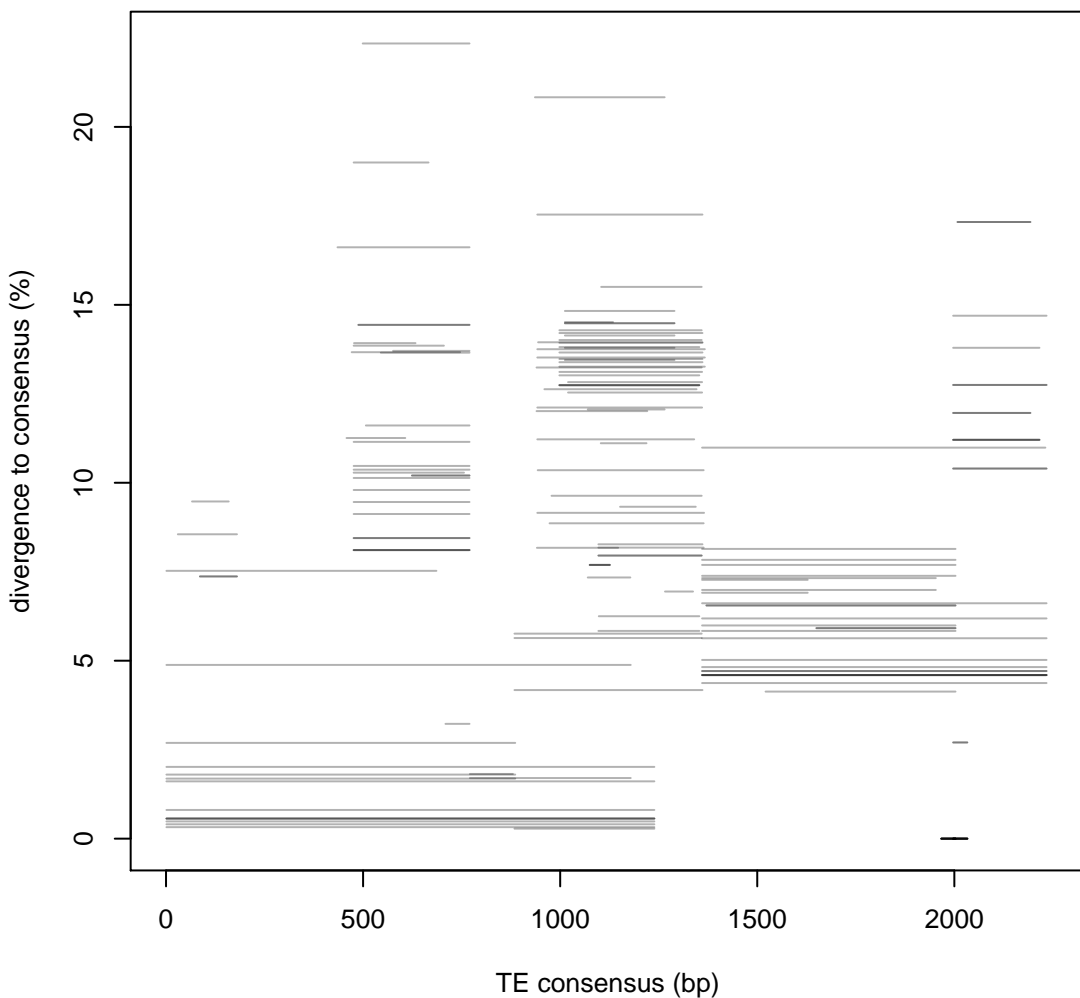
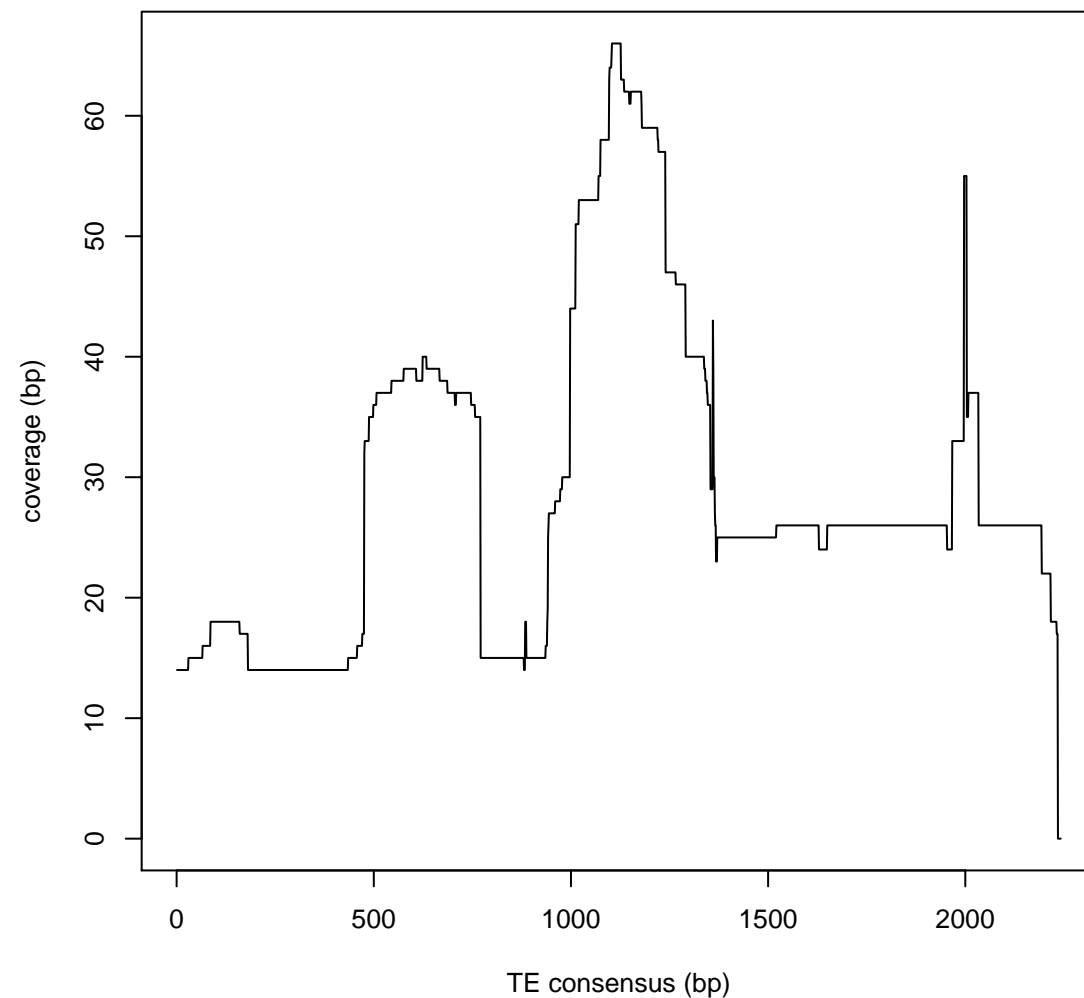


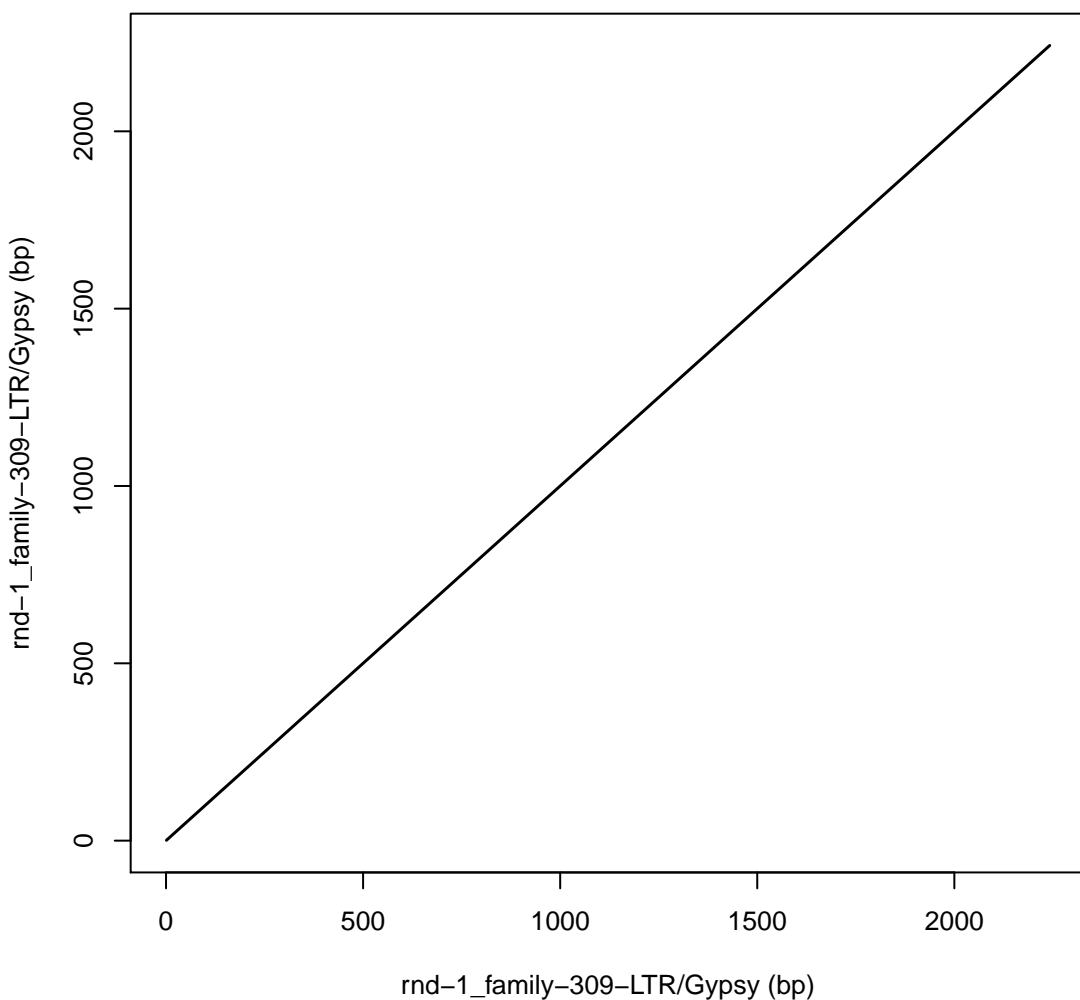
TE: rnd-1\_family-309-LTR/Gypsy  
consensus size: 2242bp; fragments: 166; full length: 0 (>=2017.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

