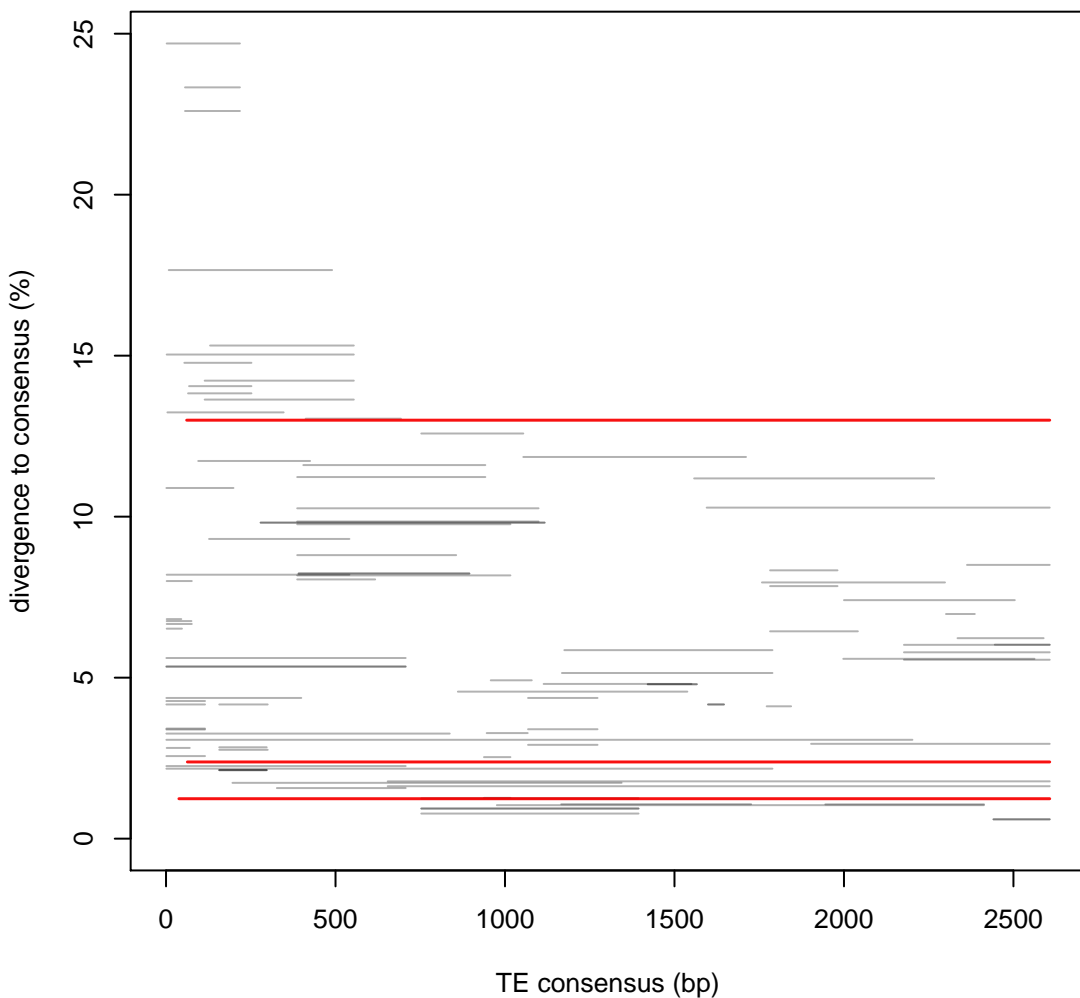
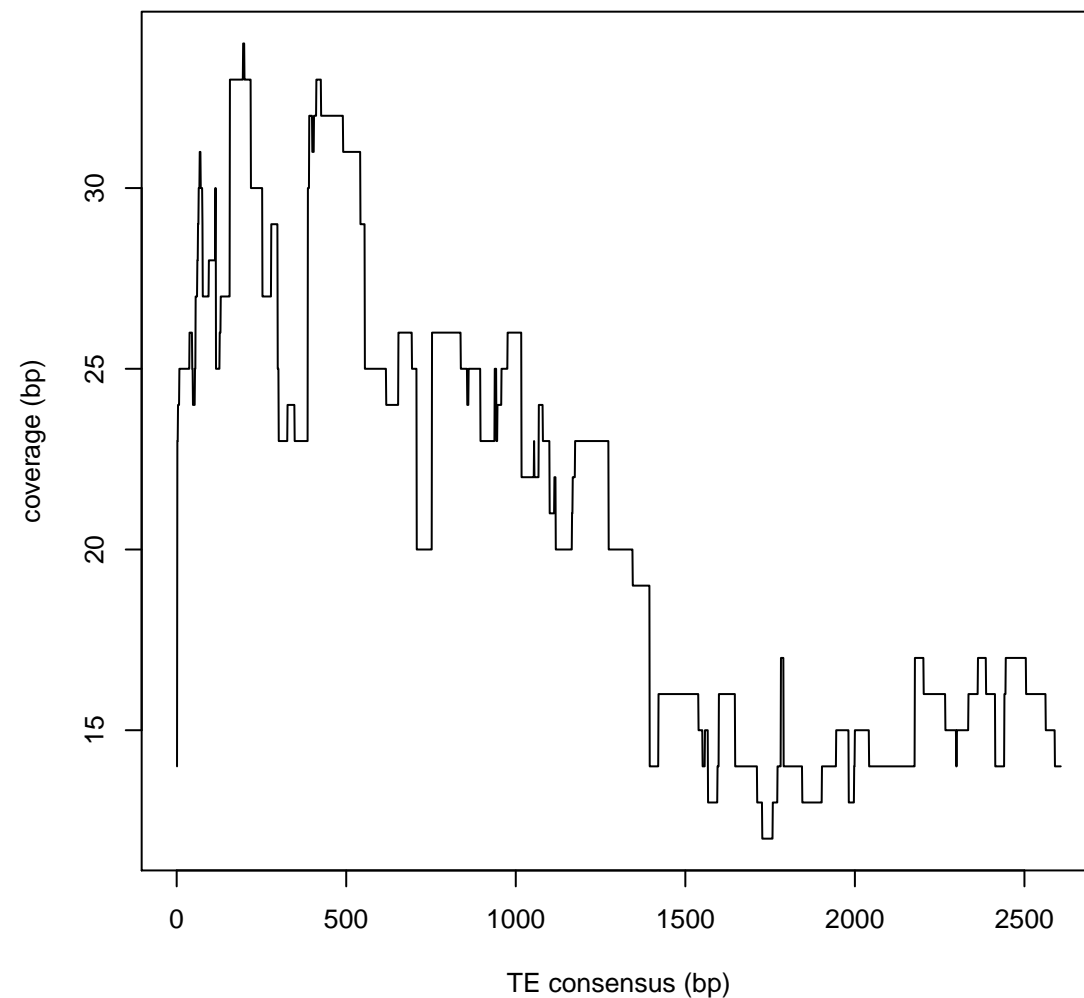


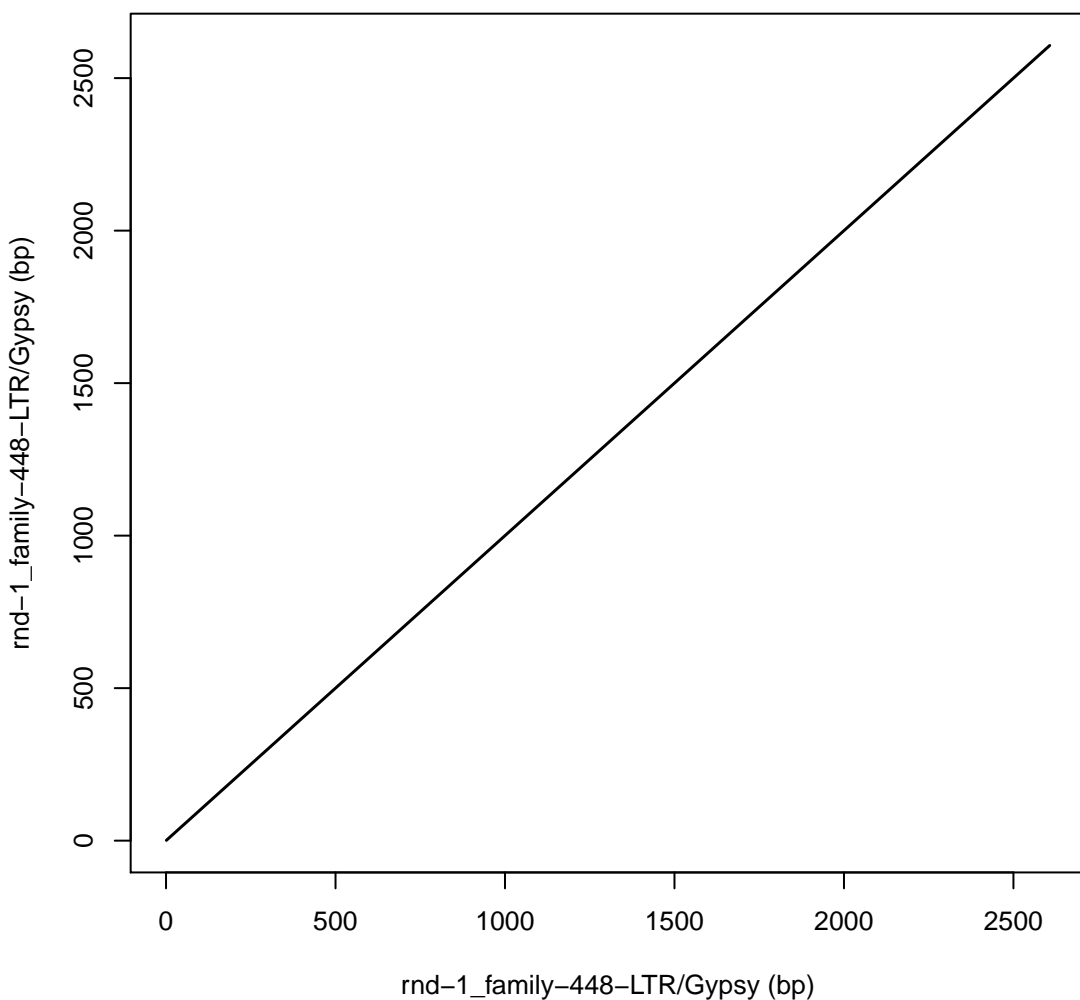
TE: rnd-1\_family-448-LTR/Gypsy  
consensus size: 2607bp; fragments: 105; full length: 3 (>=2346.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

