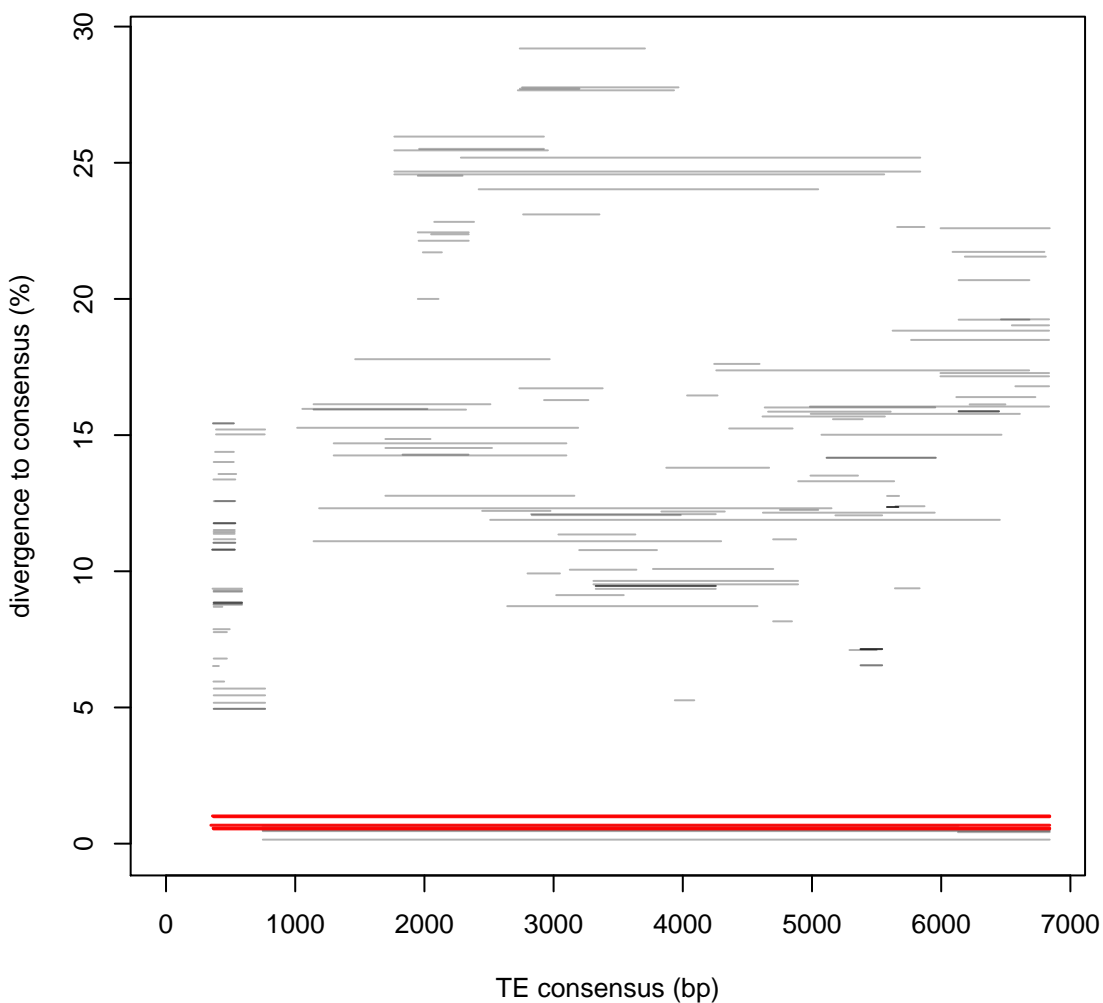
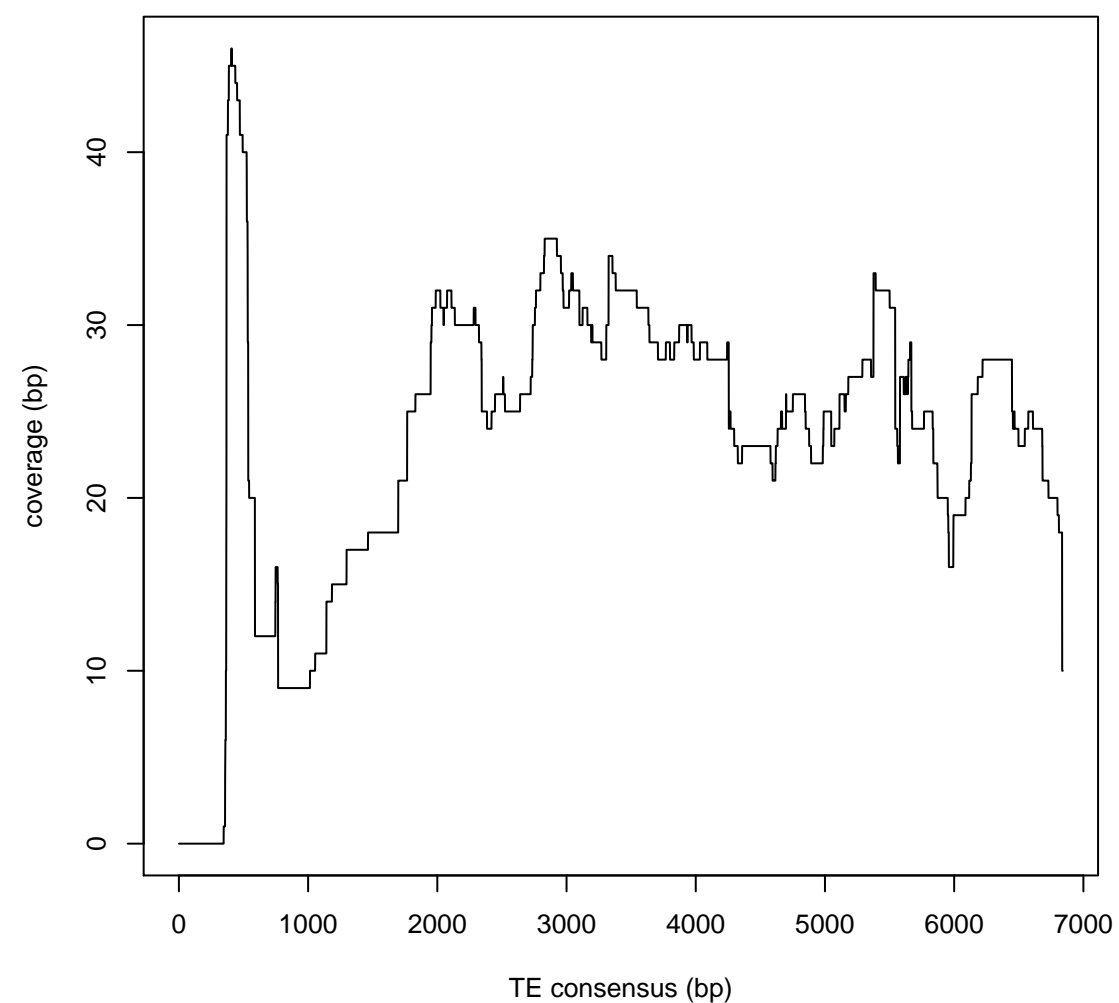


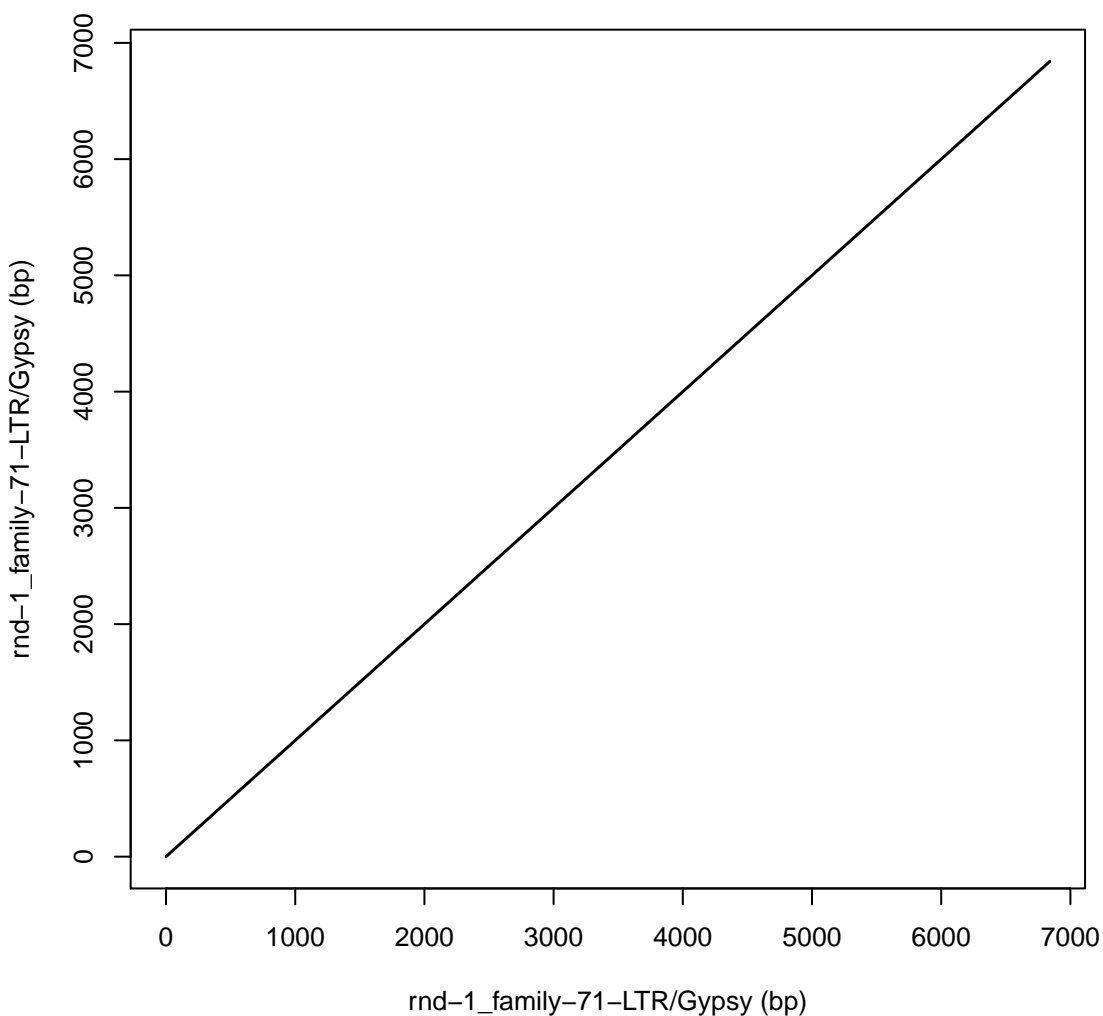
TE: rnd-1\_family-71-LTR/Gypsy  
consensus size: 6840bp; fragments: 157; full length: 5 (>=6156bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

