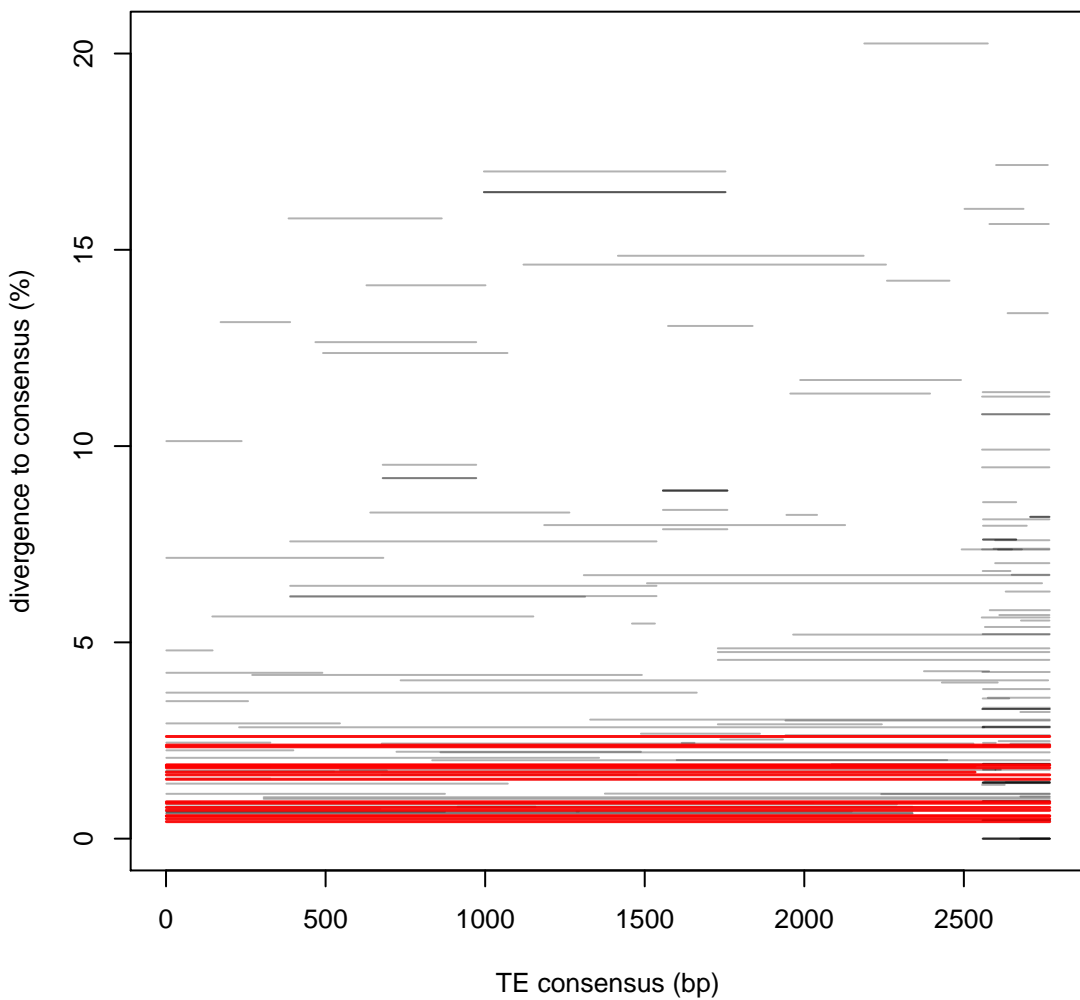
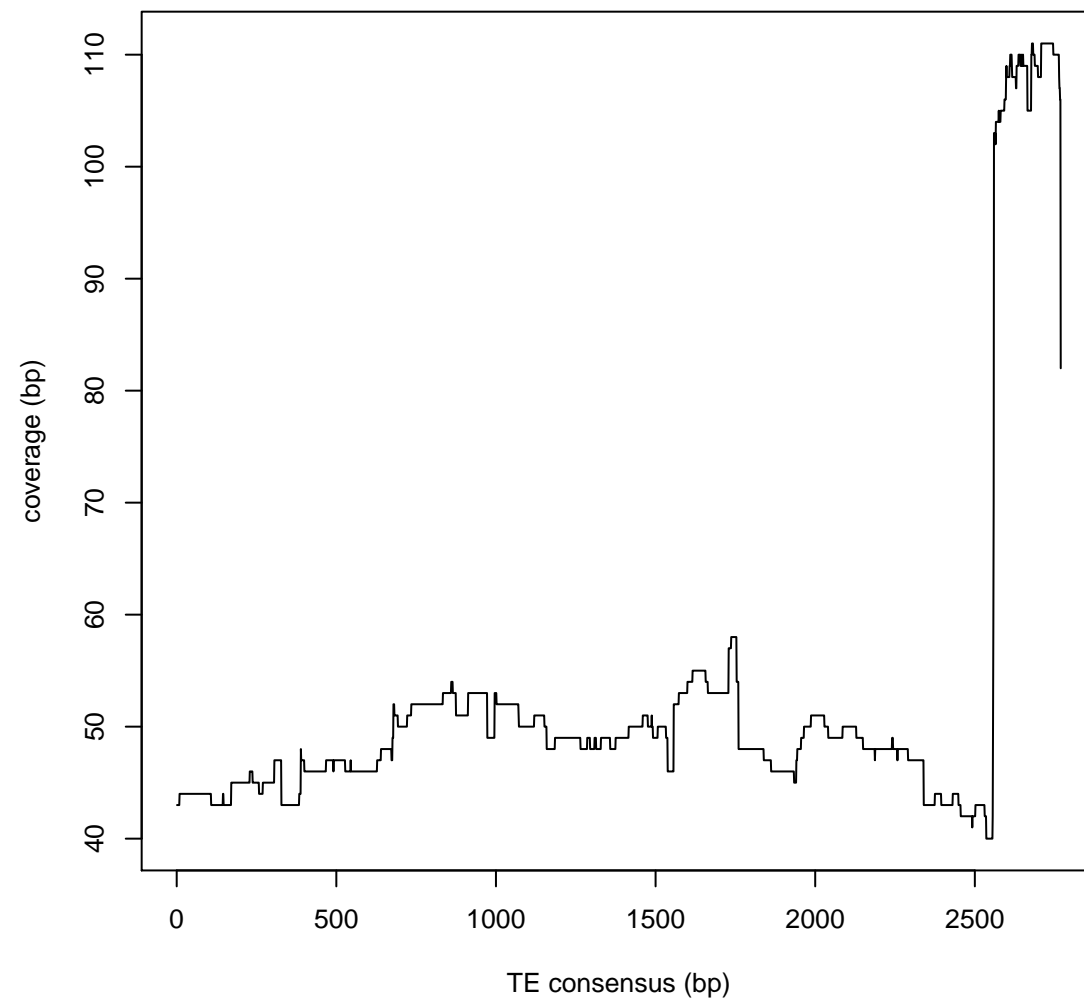


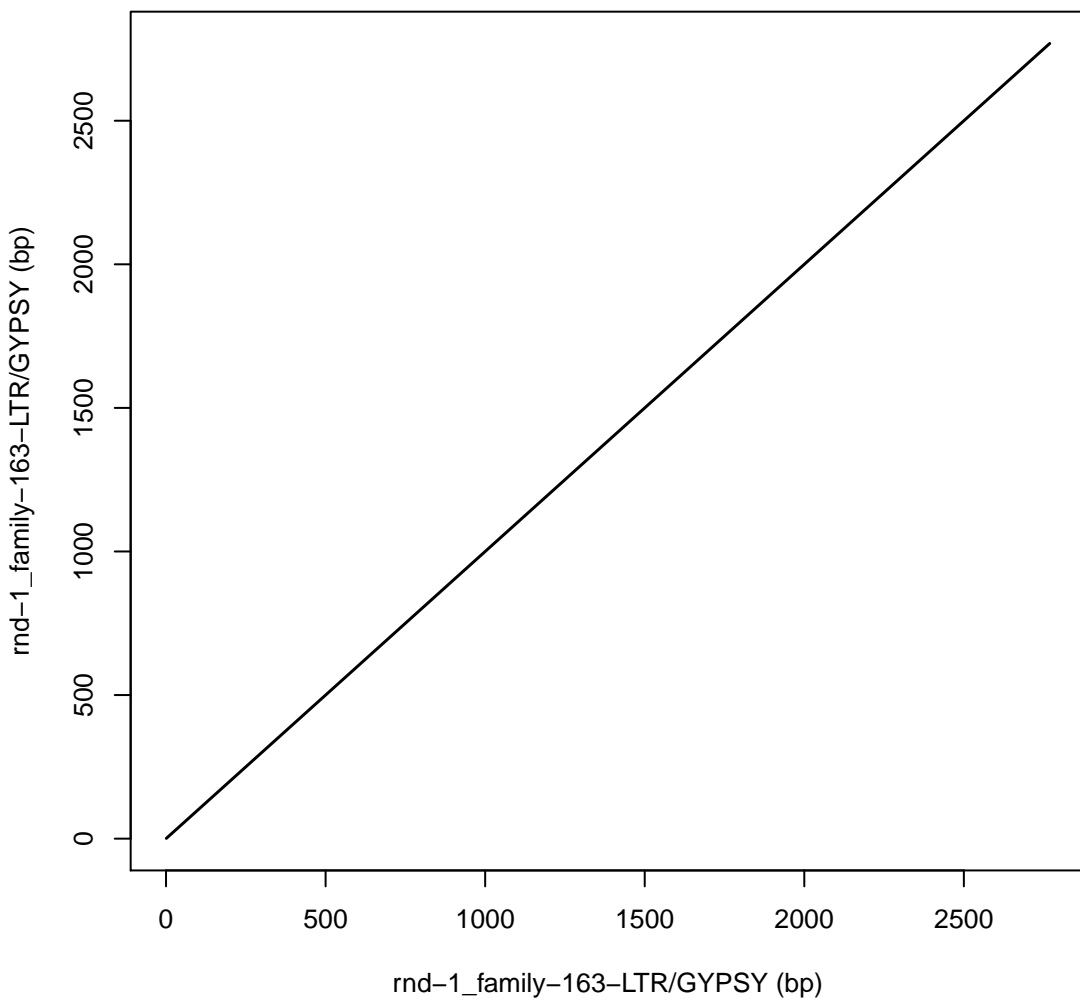
TE: rnd-1\_family-163-LTR/GYPSY  
consensus size: 2769bp; fragments: 204; full length: 18 (>=2492.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

