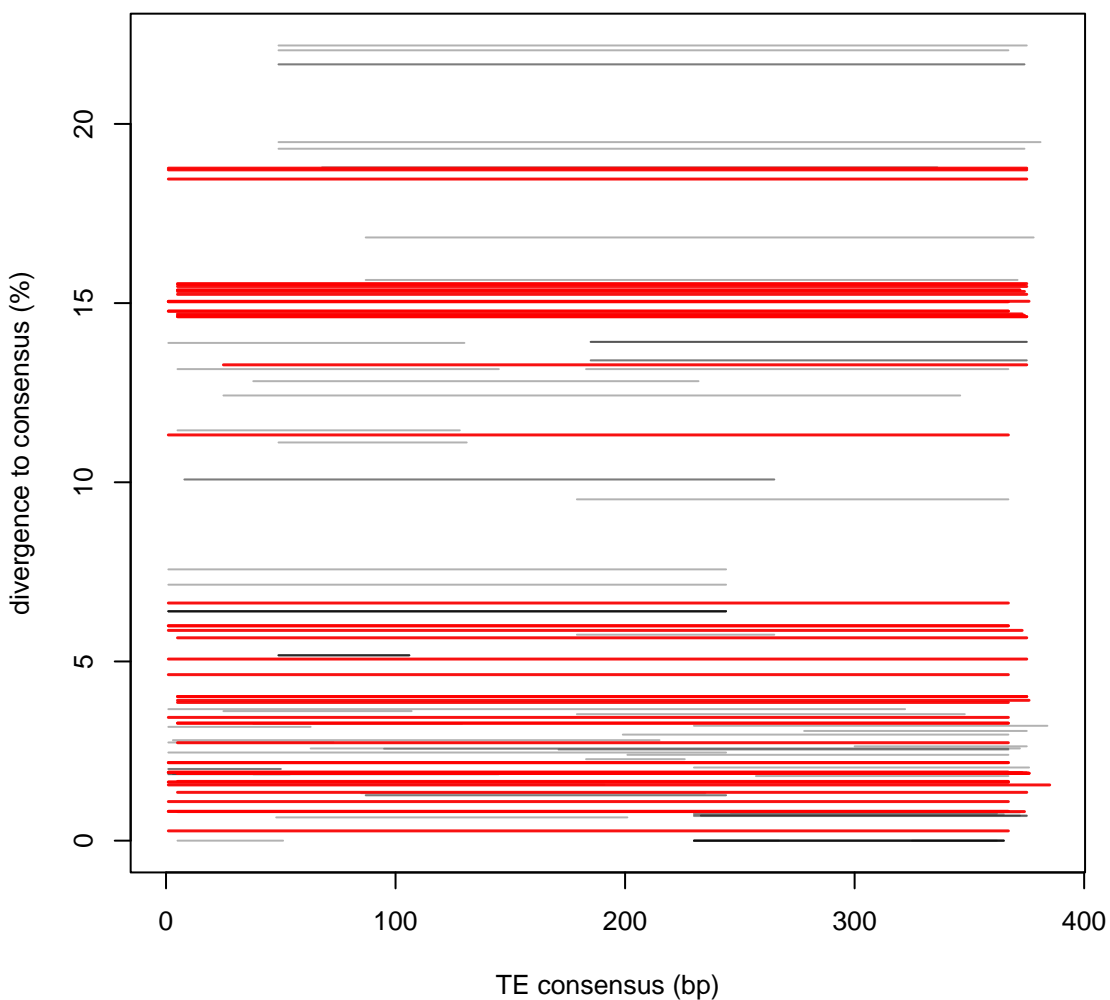
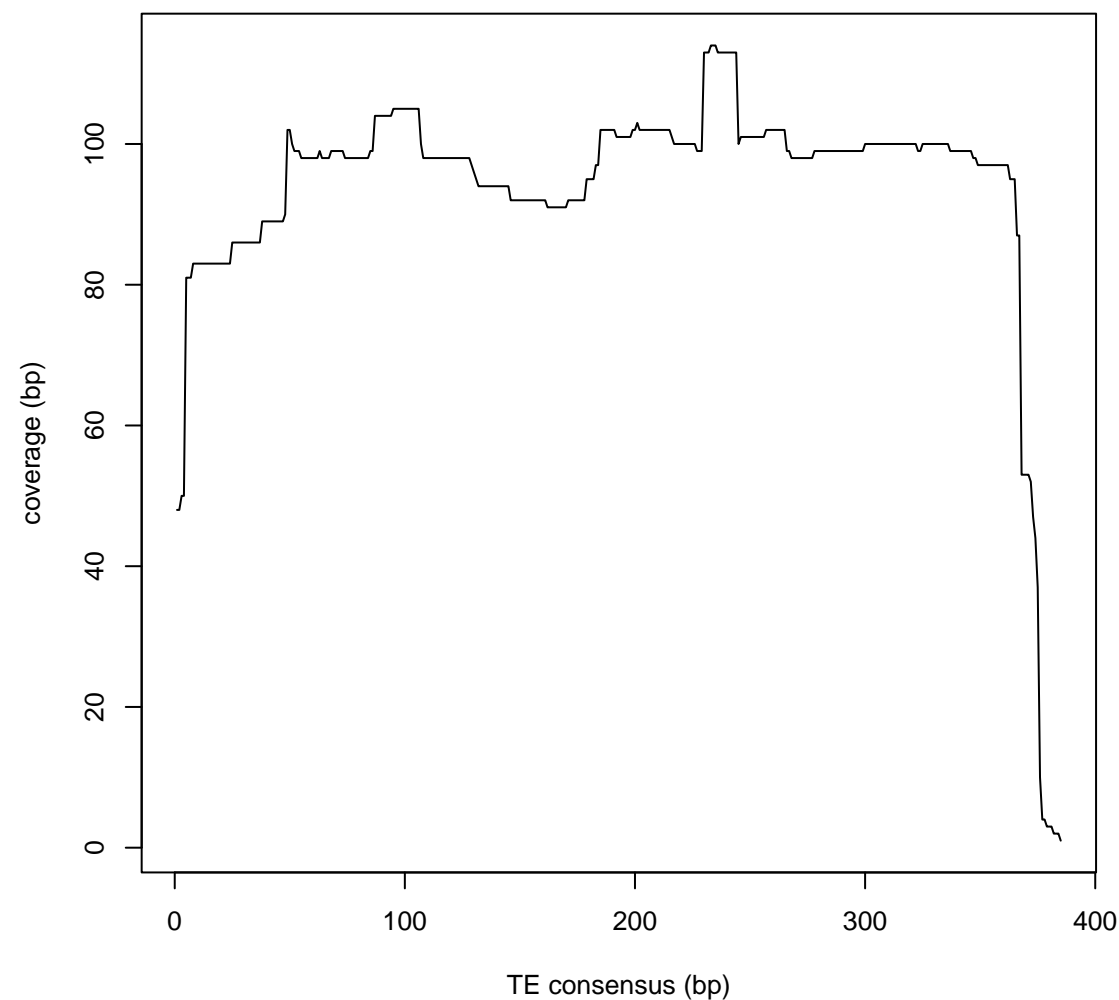


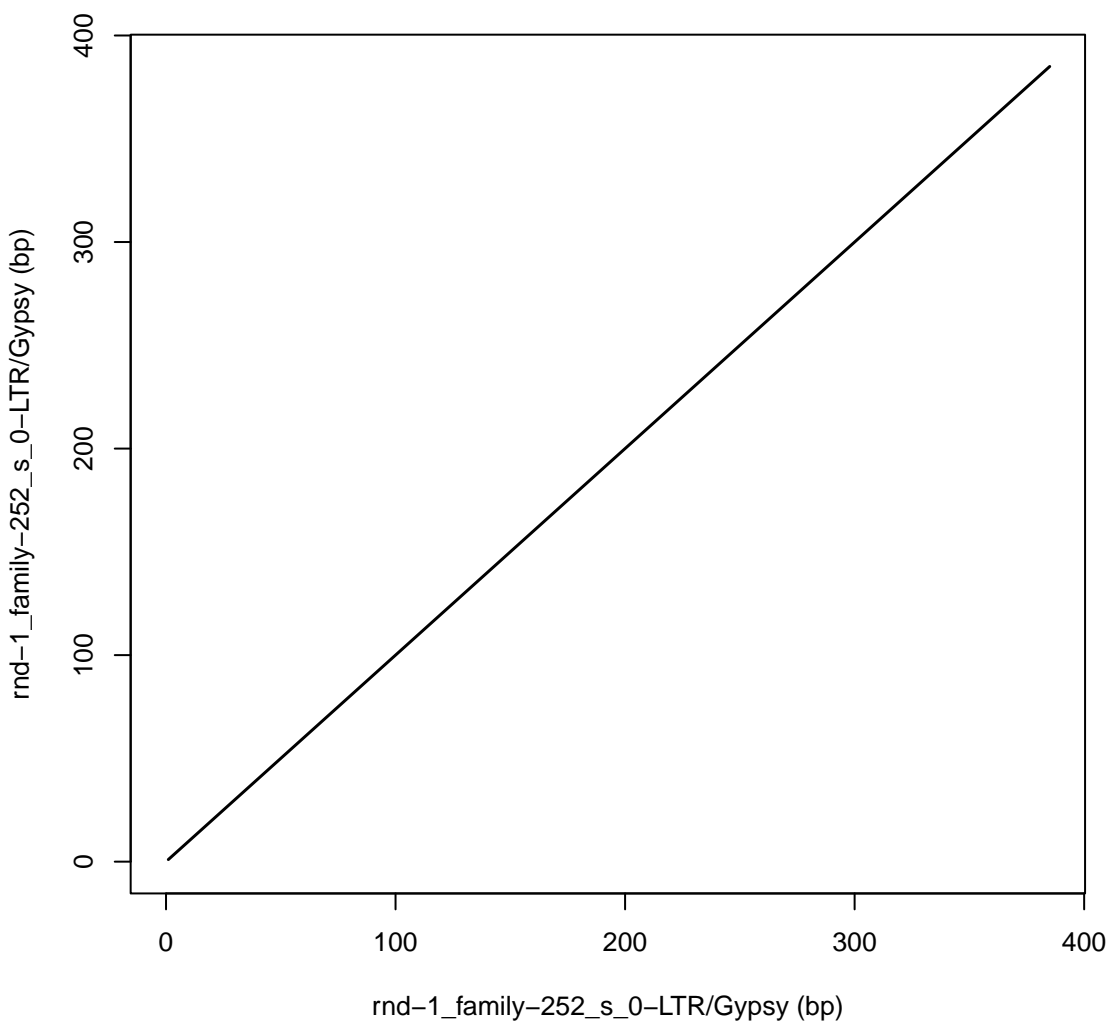
TE: rnd-1\_family-252\_s\_0-LTR/Gypsy  
consensus size: 385bp; fragments: 145; full length: 57 (>=346.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

