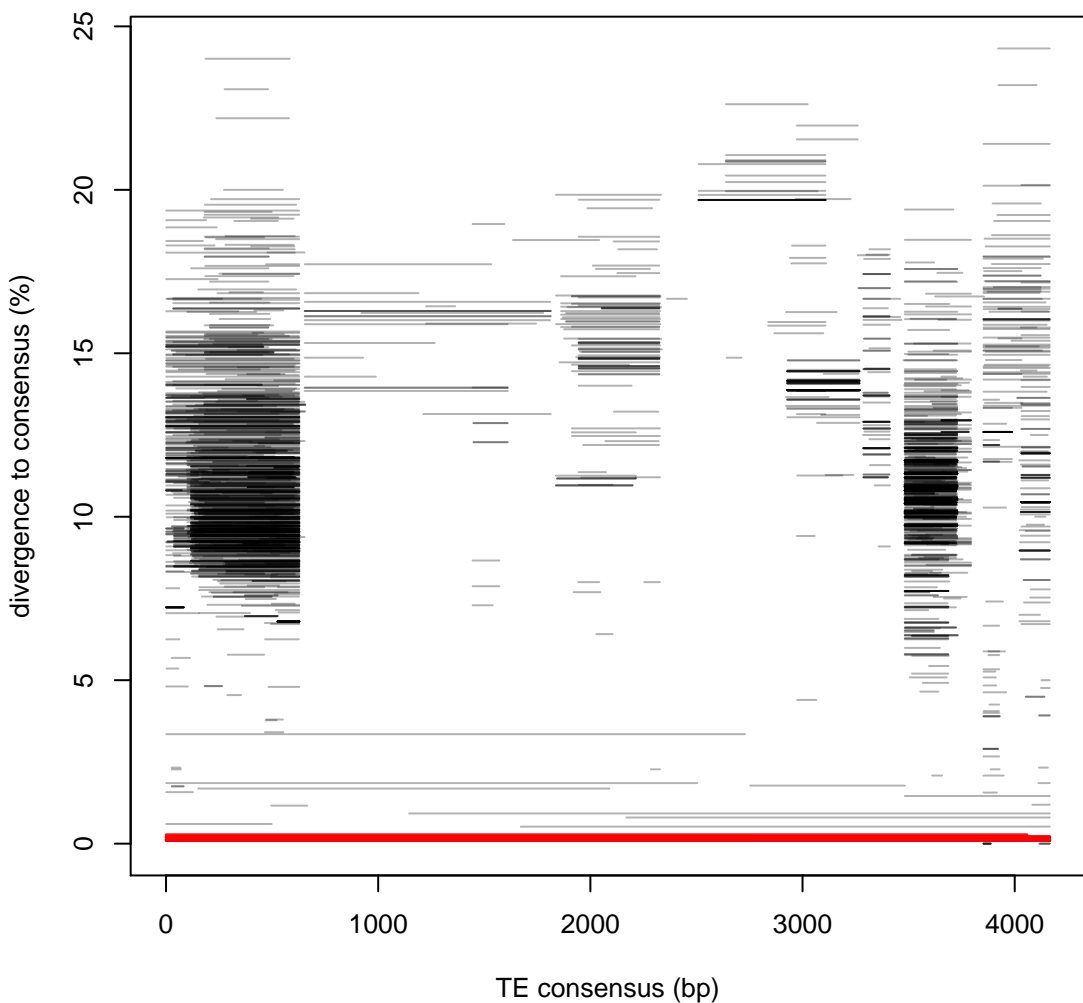
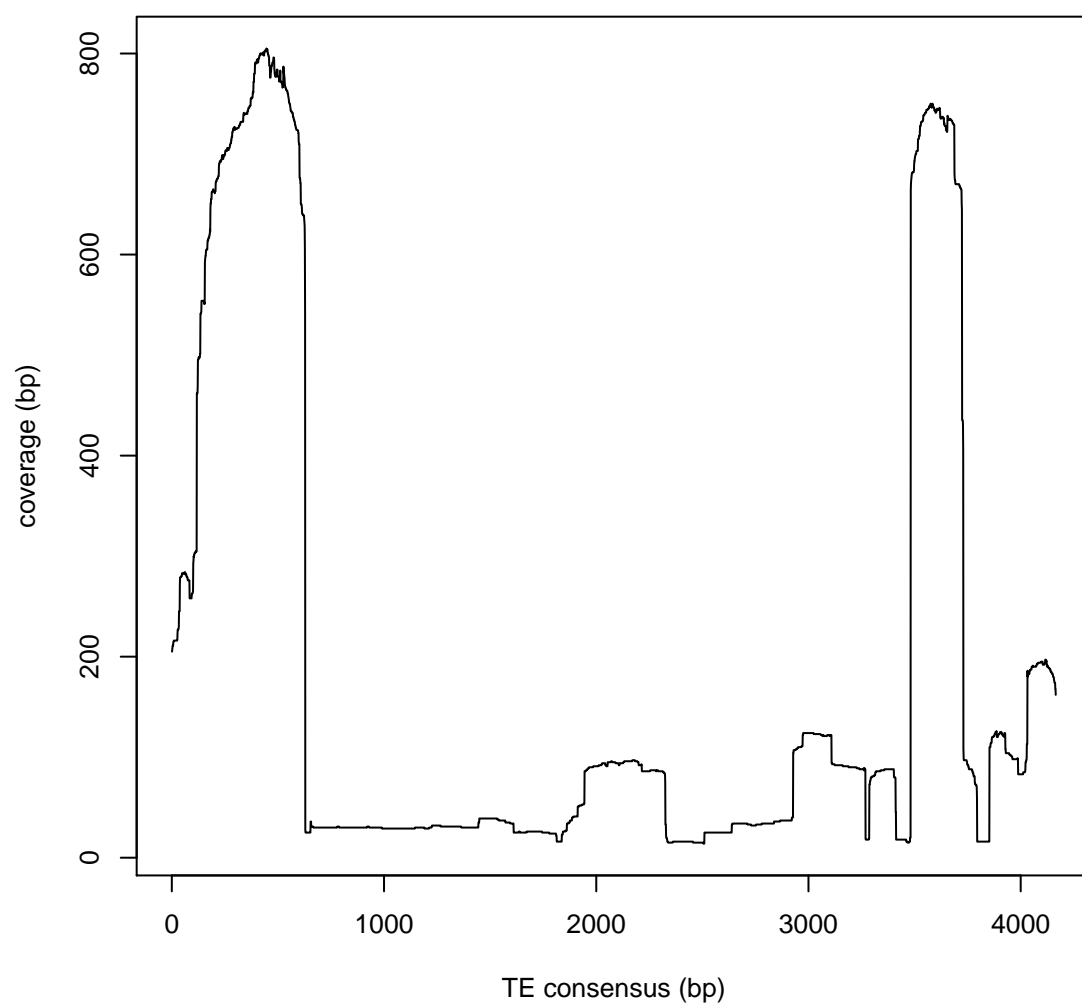


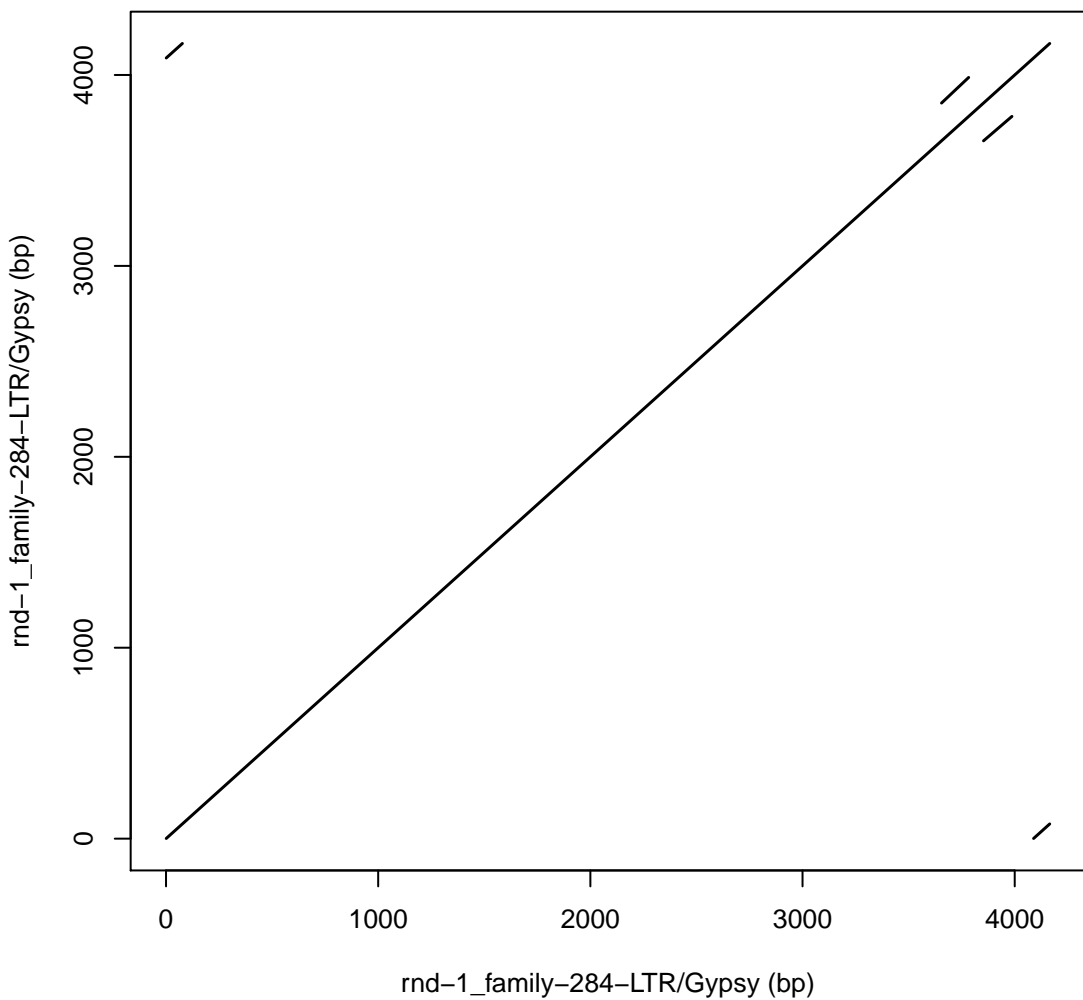
TE: rnd-1_family-284-LTR/Gypsy
consensus size: 4165bp; fragments: 2306; full length: 9 (>=3748.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

