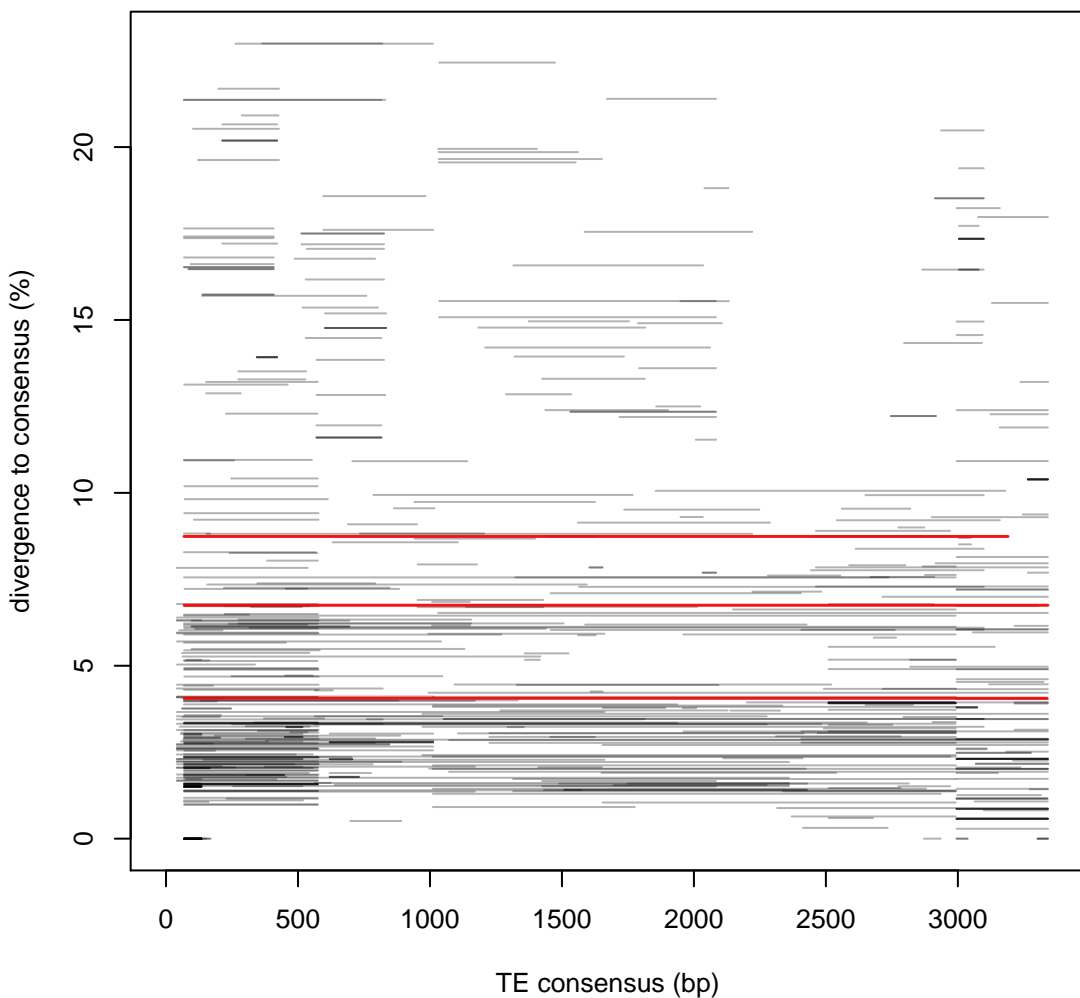
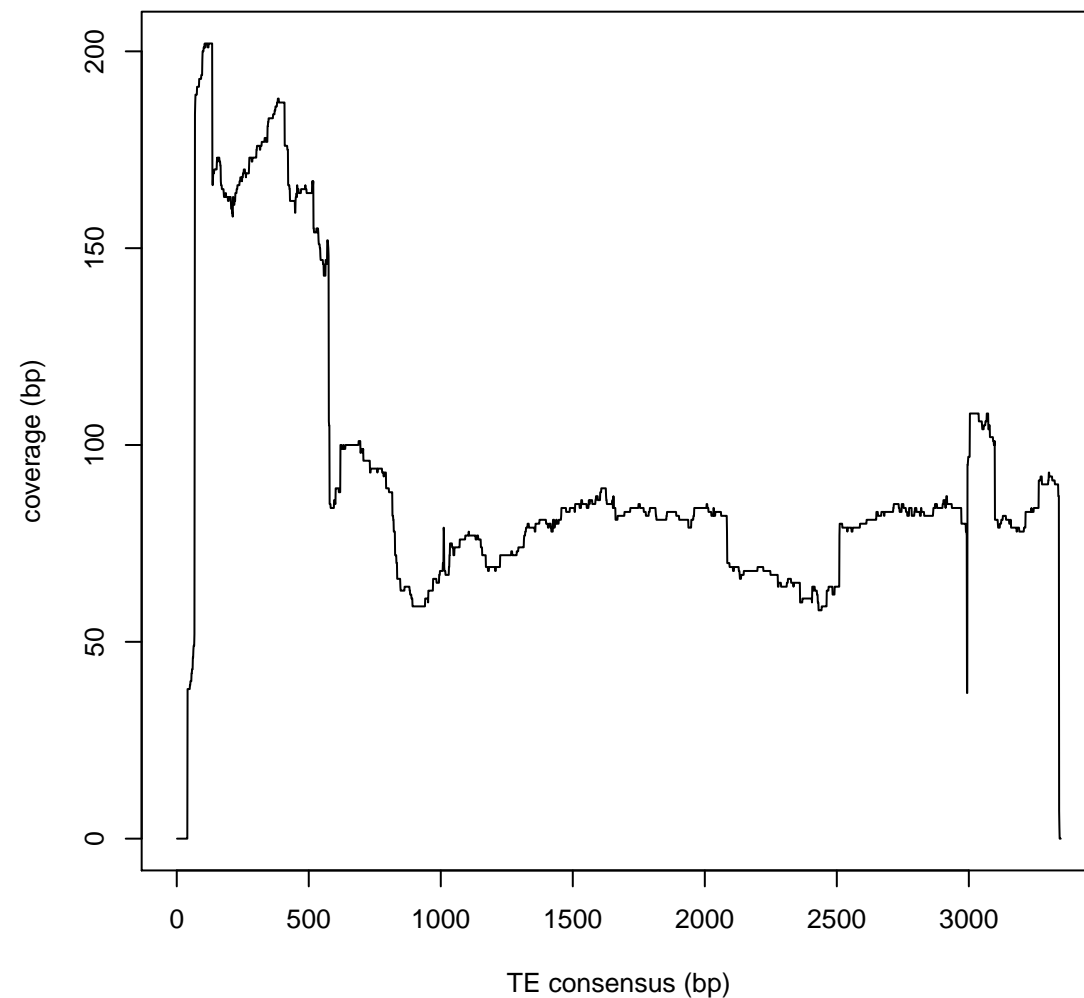


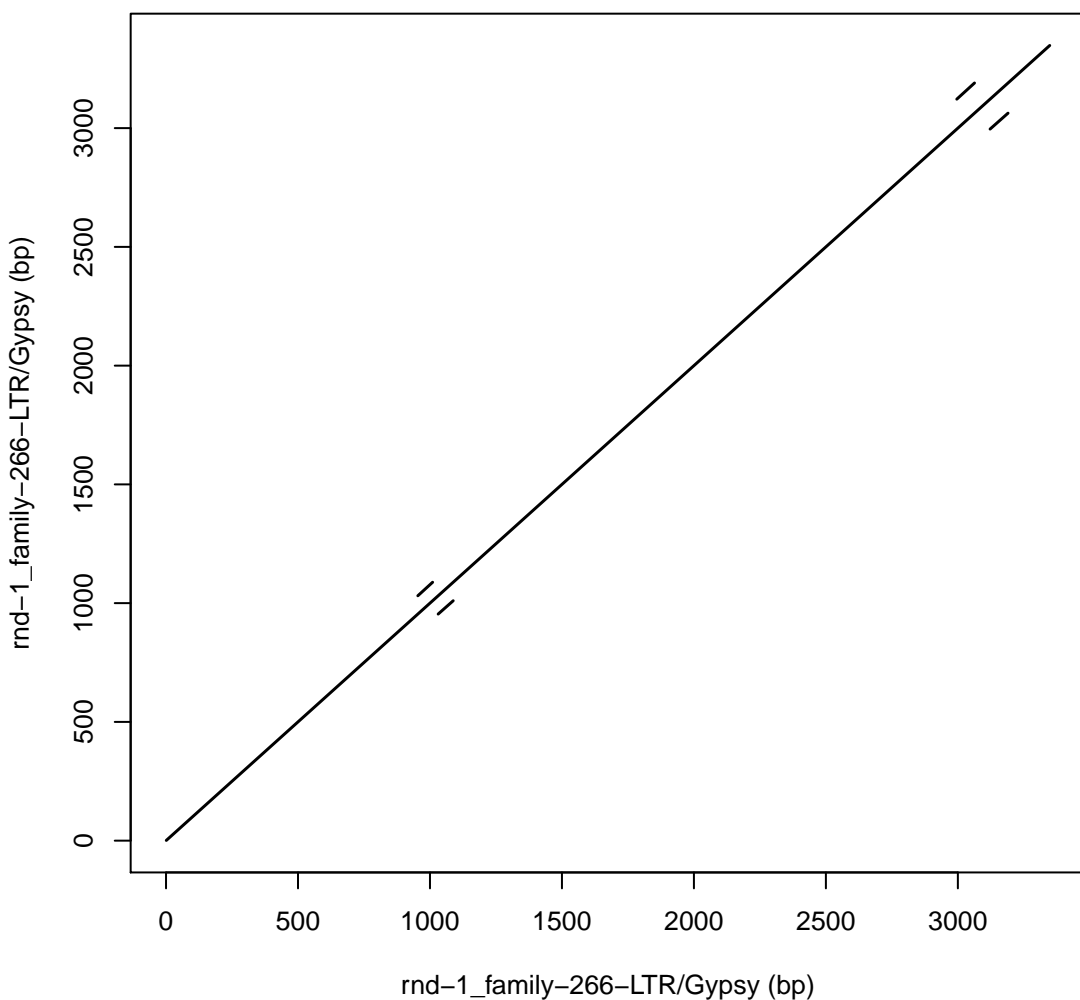
TE: rnd-1_family-266-LTR/Gypsy
consensus size: 3348bp; fragments: 575; full length: 3 (>=3013.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

