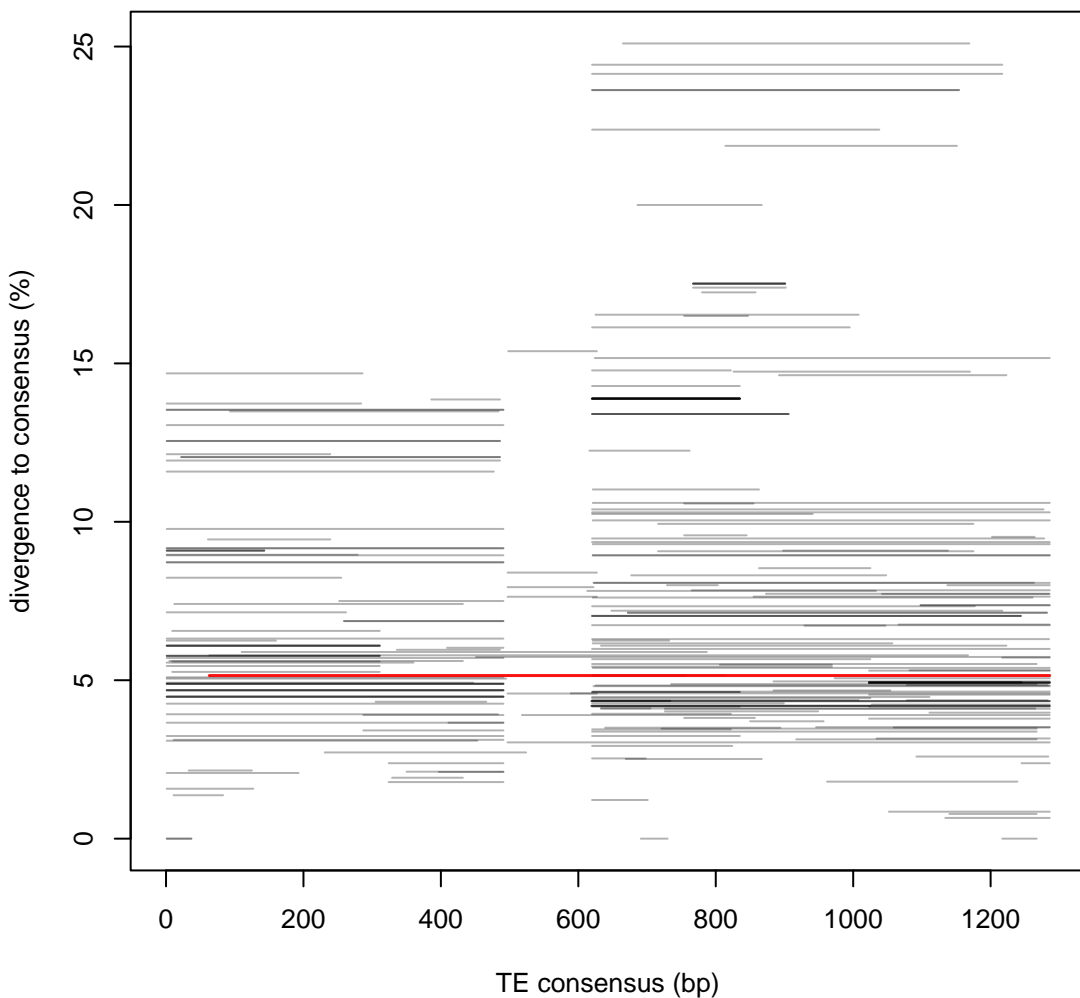
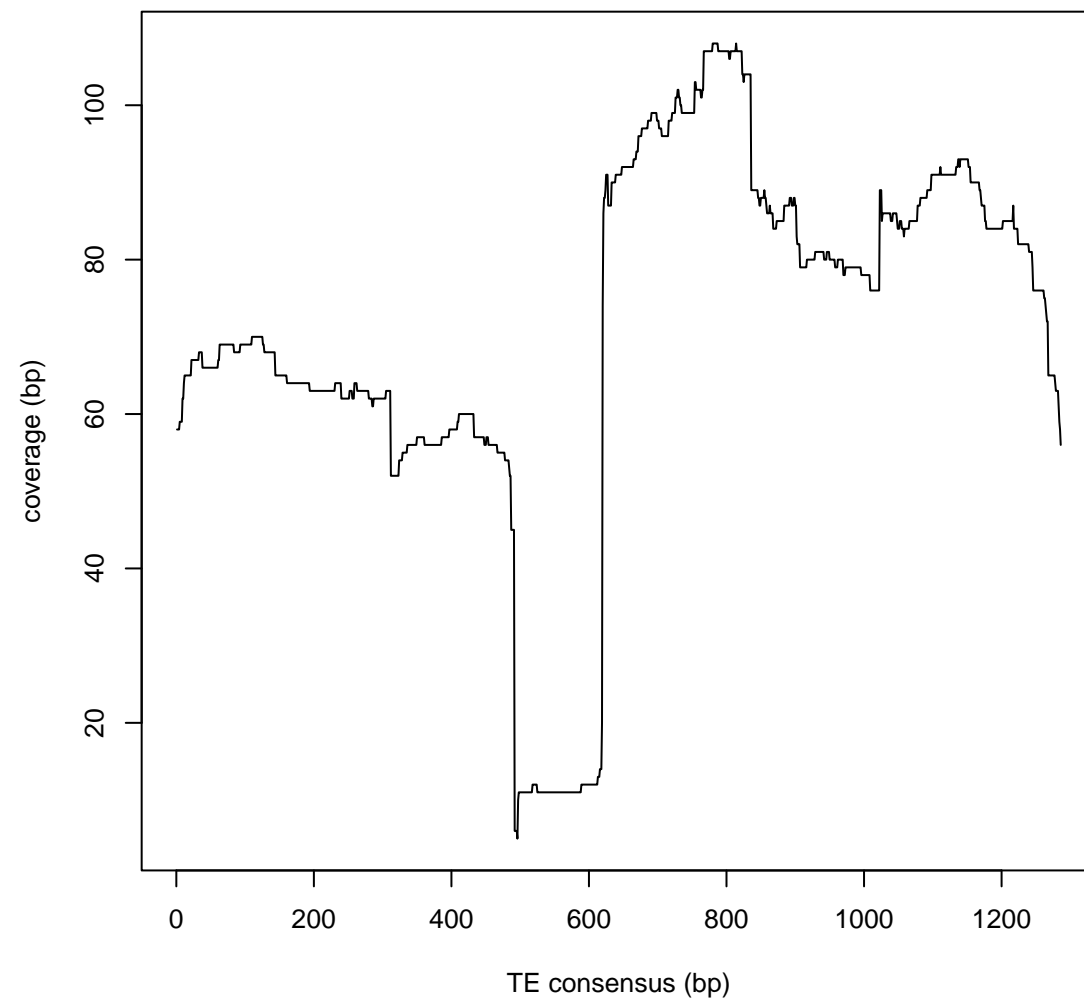


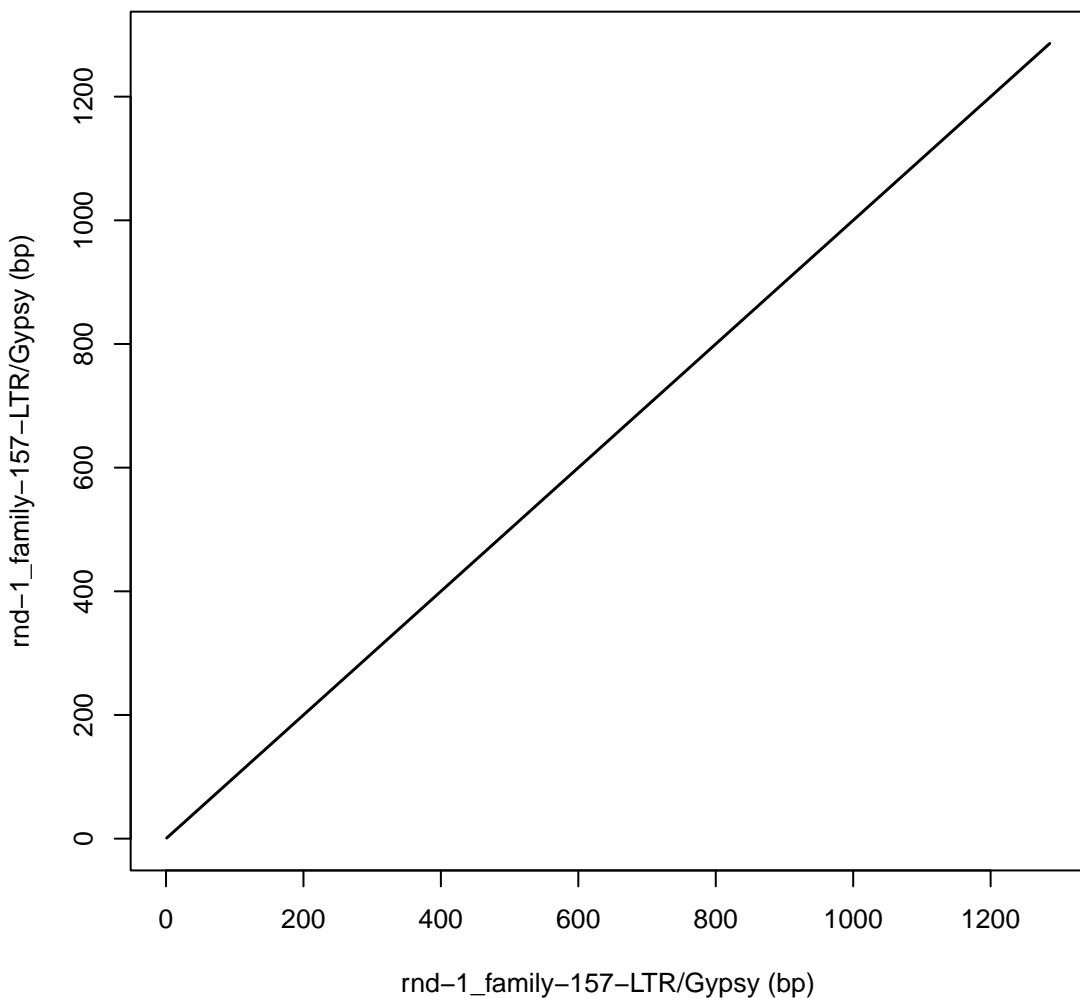
TE: rnd-1_family-157-LTR/Gypsy
consensus size: 1286bp; fragments: 257; full length: 1 (>=1157.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

