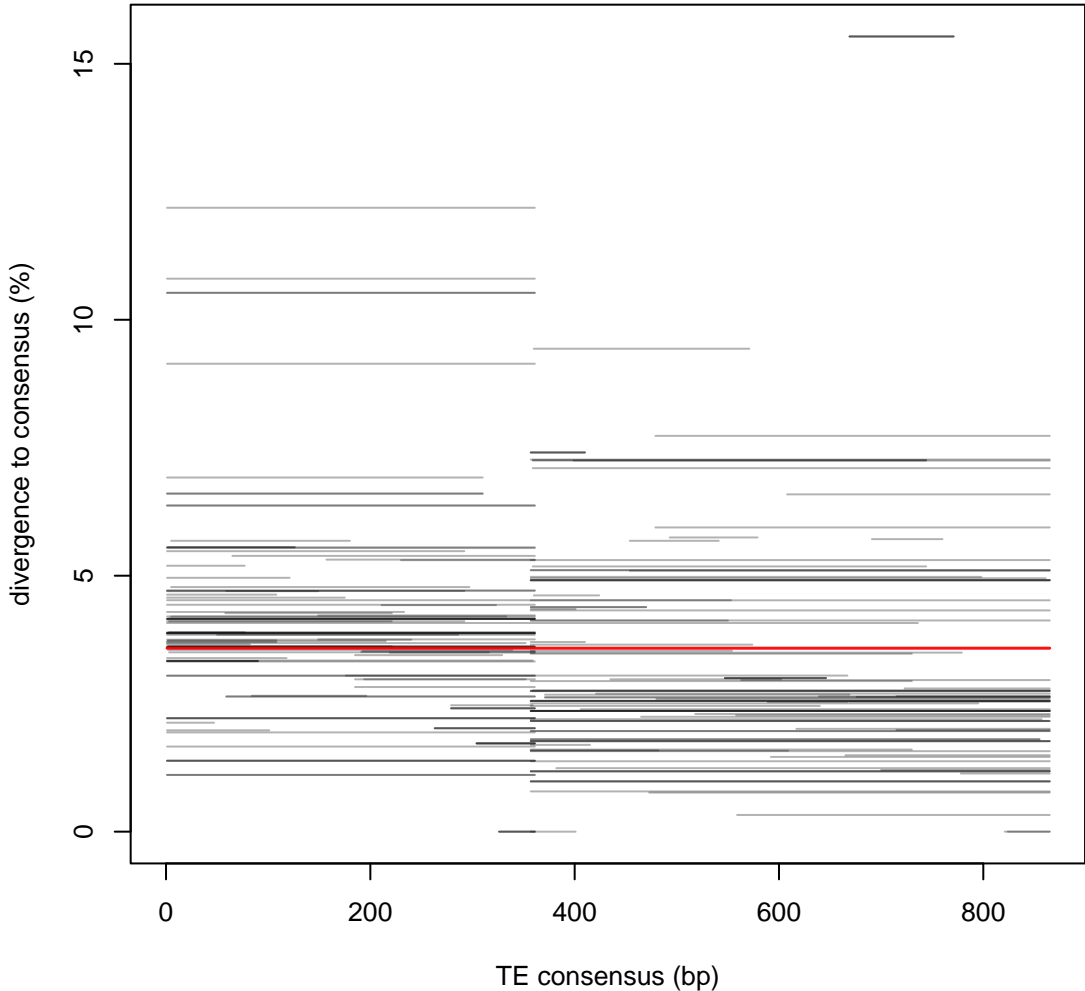
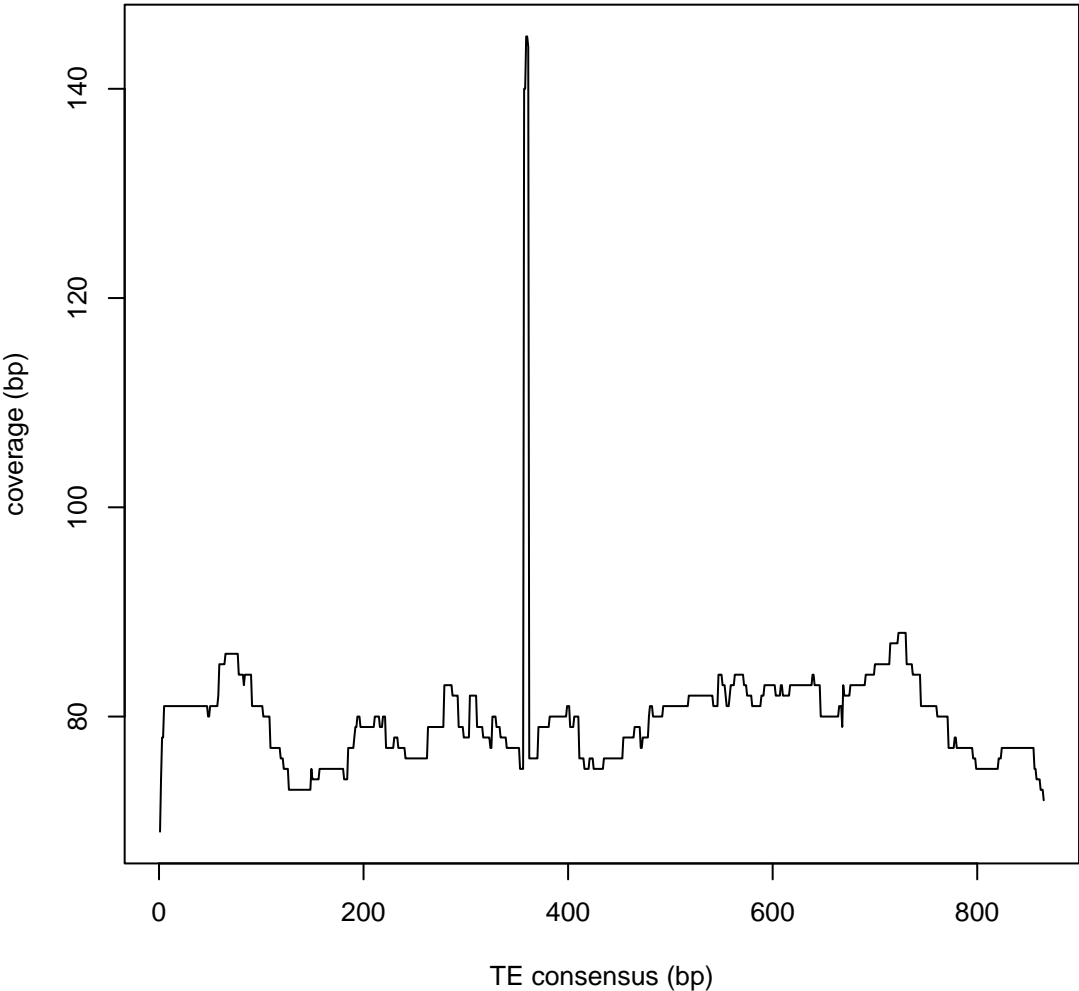


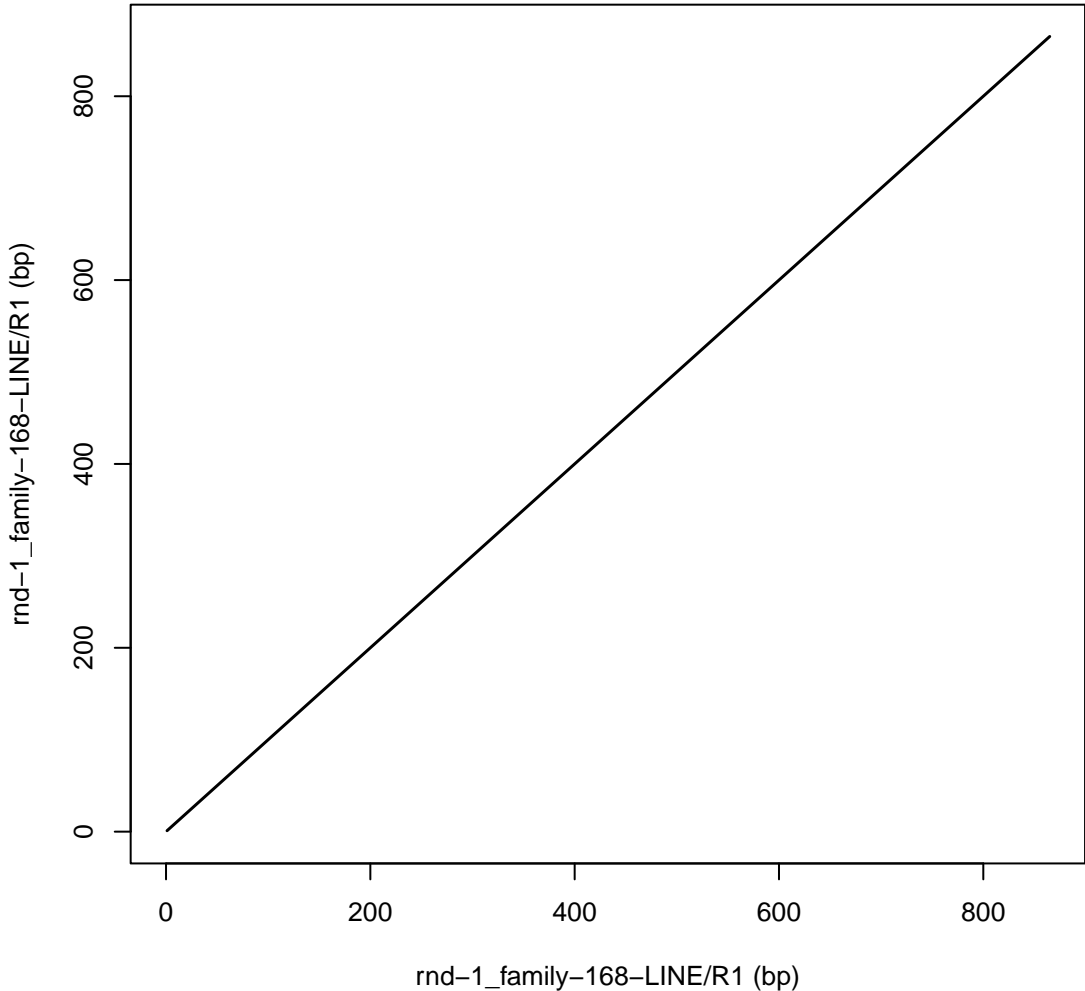
TE: rnd-1_family-168-LINE/R1
consensus size: 865bp; fragments: 229; full length: 1 (>=778.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

