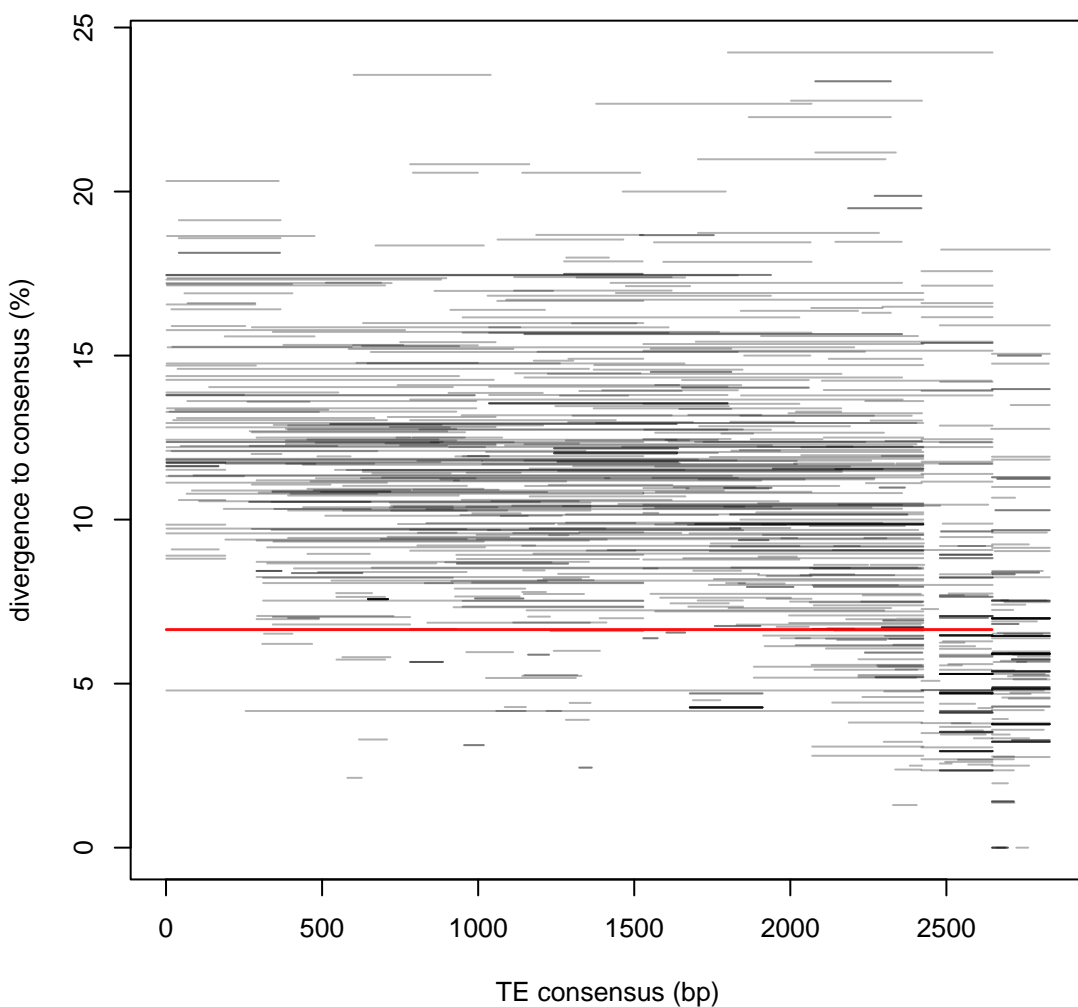
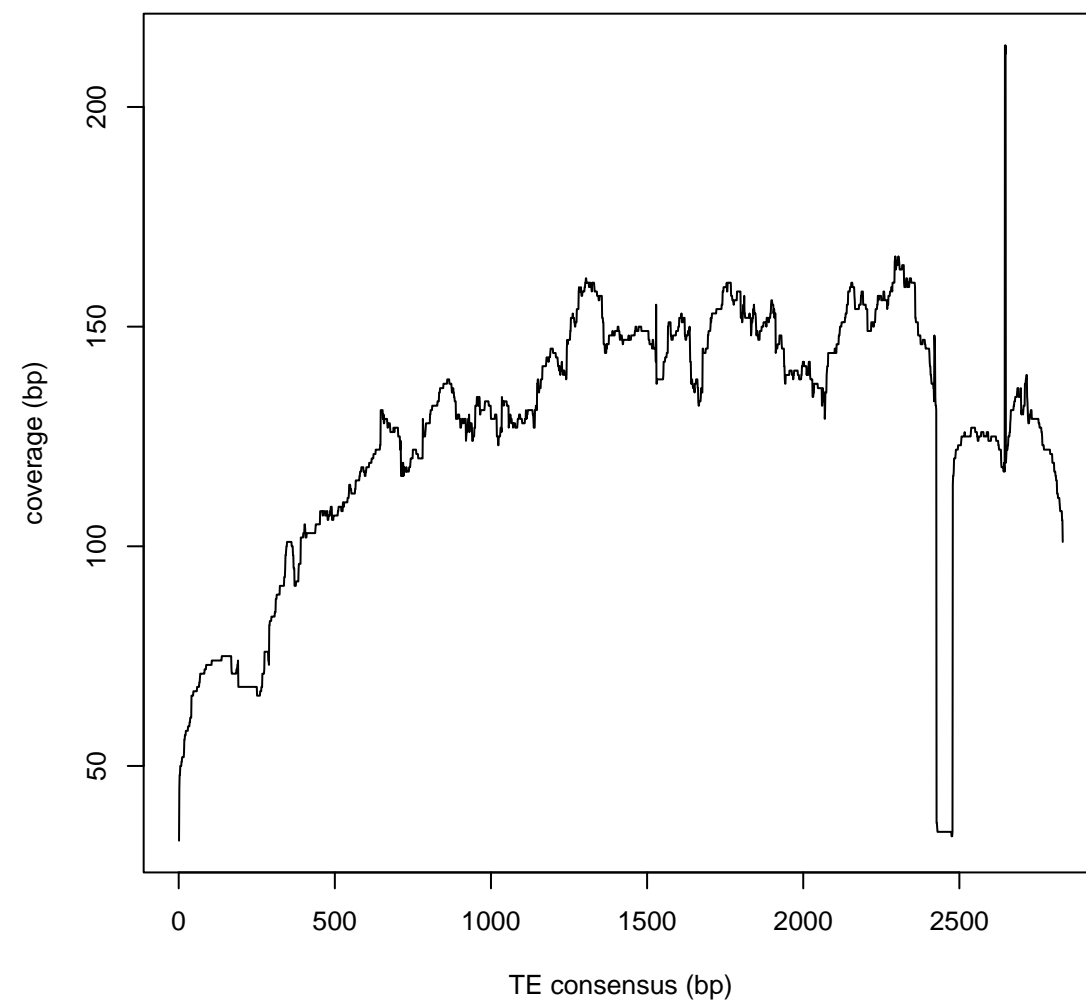


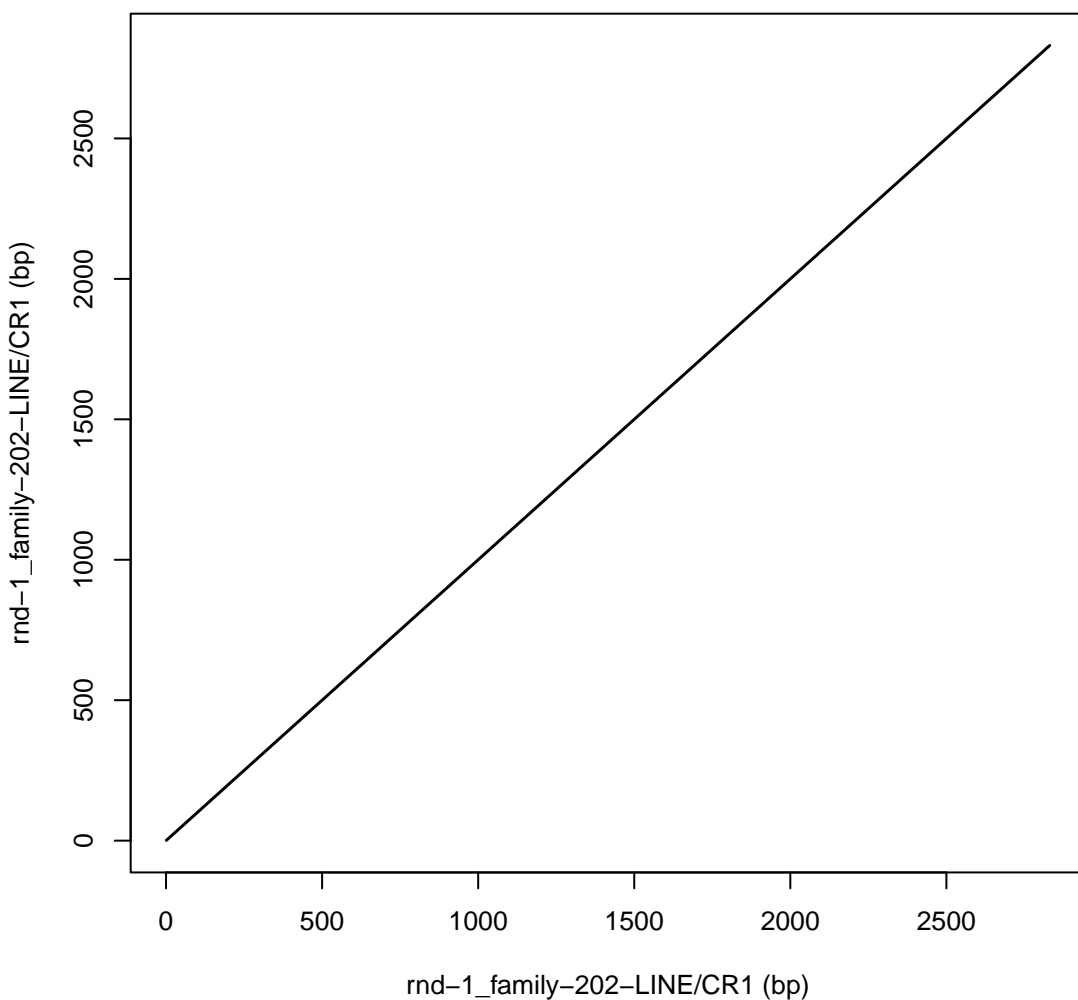
TE: rnd-1_family-202-LINE/CR1
consensus size: 2831bp; fragments: 859; full length: 1 (≥ 2547.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

