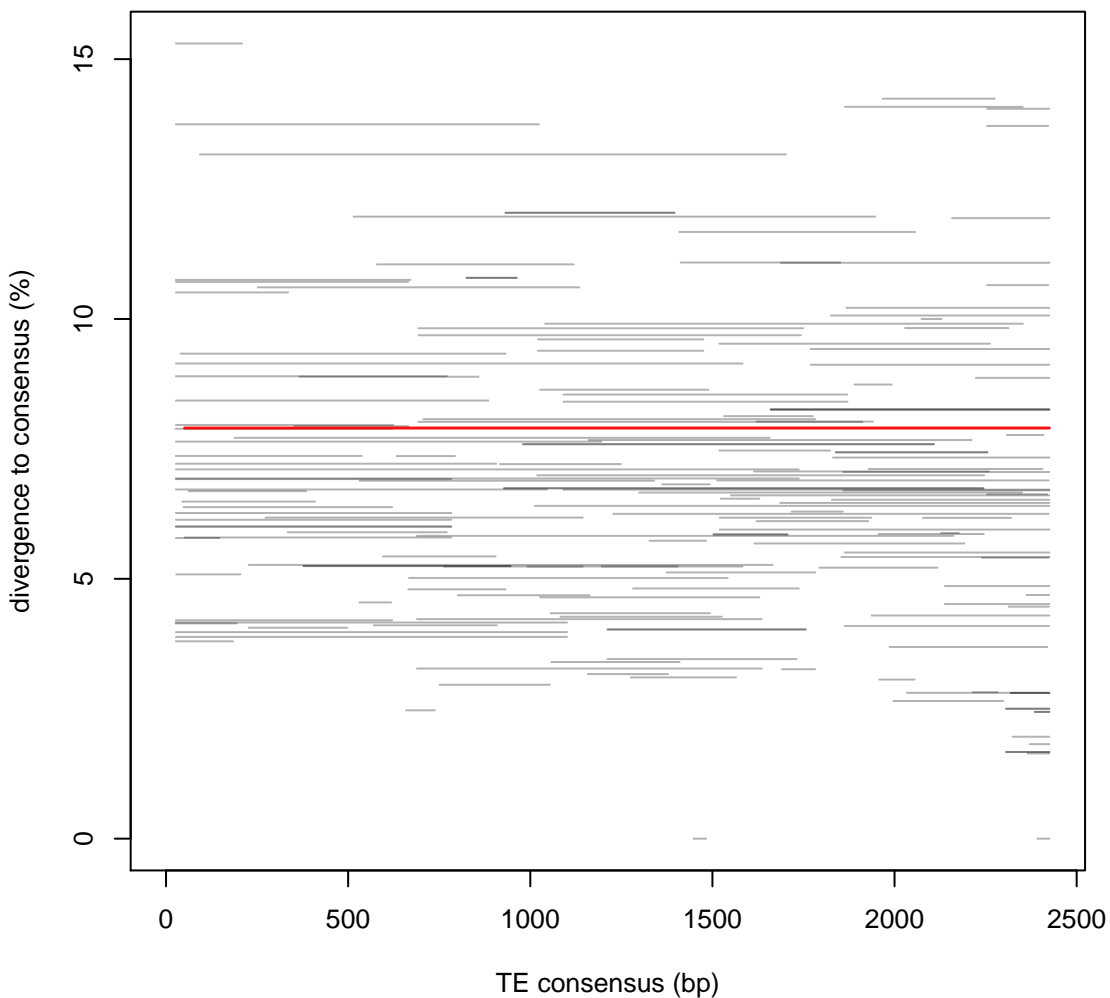
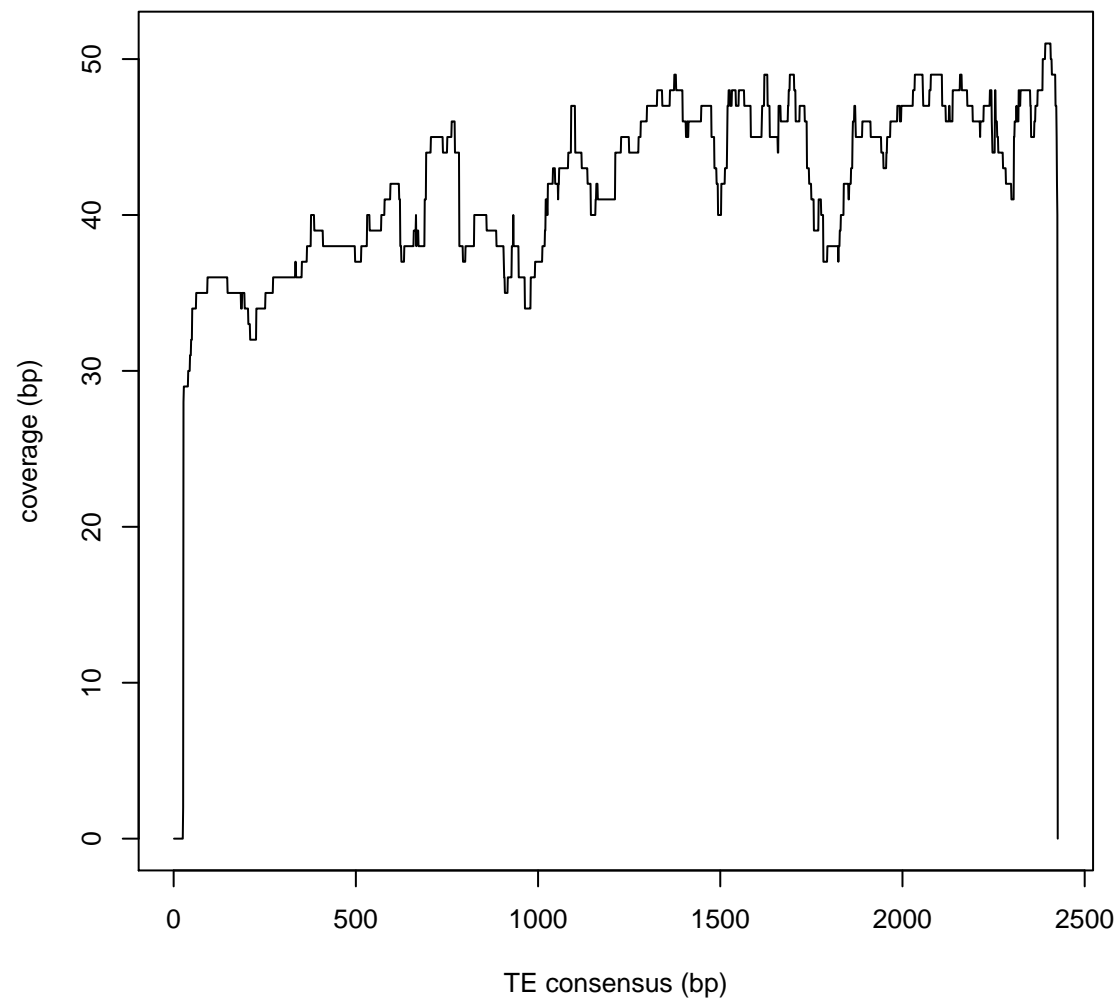


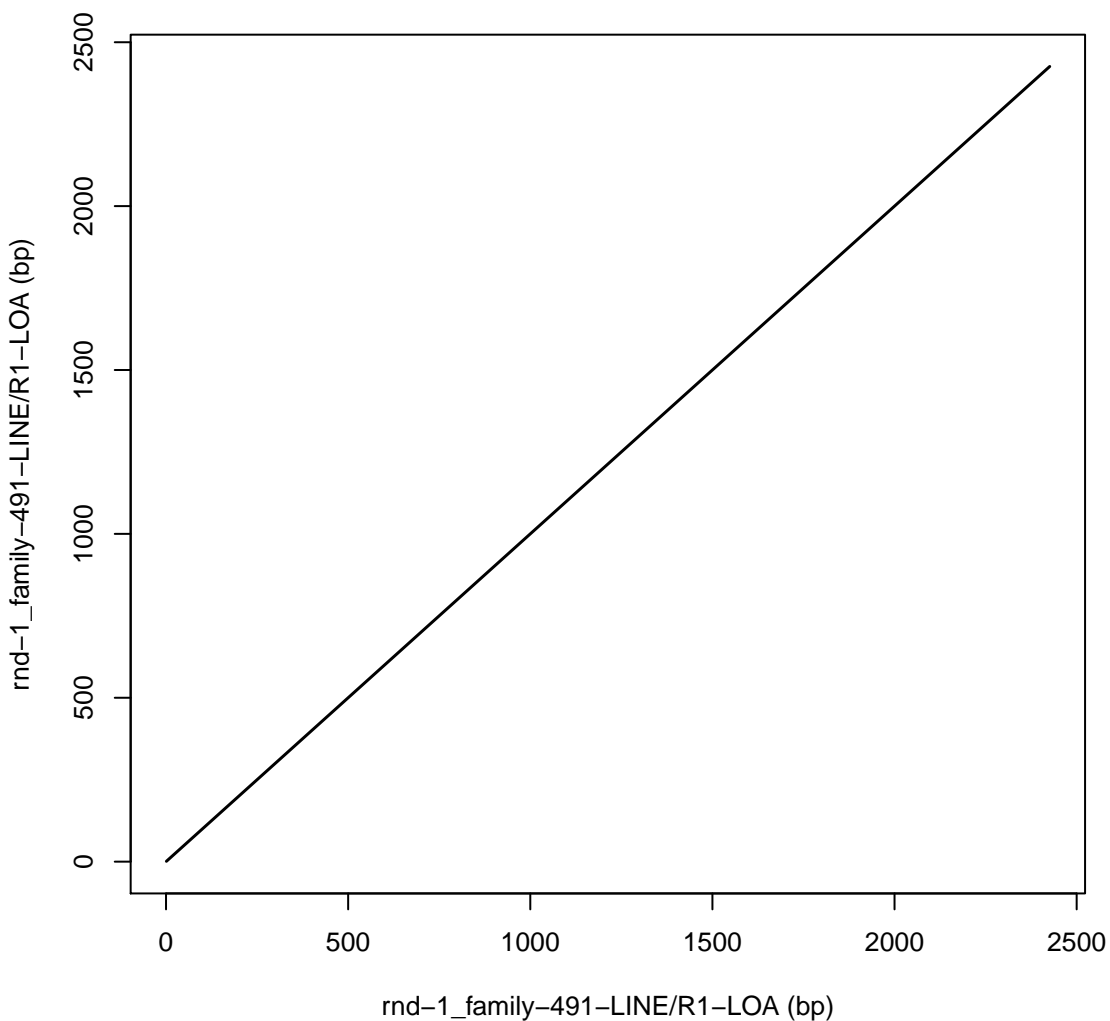
TE: rnd-1_family-491-LINE/R1-LOA
consensus size: 2426bp; fragments: 179; full length: 1 (>=2183.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

