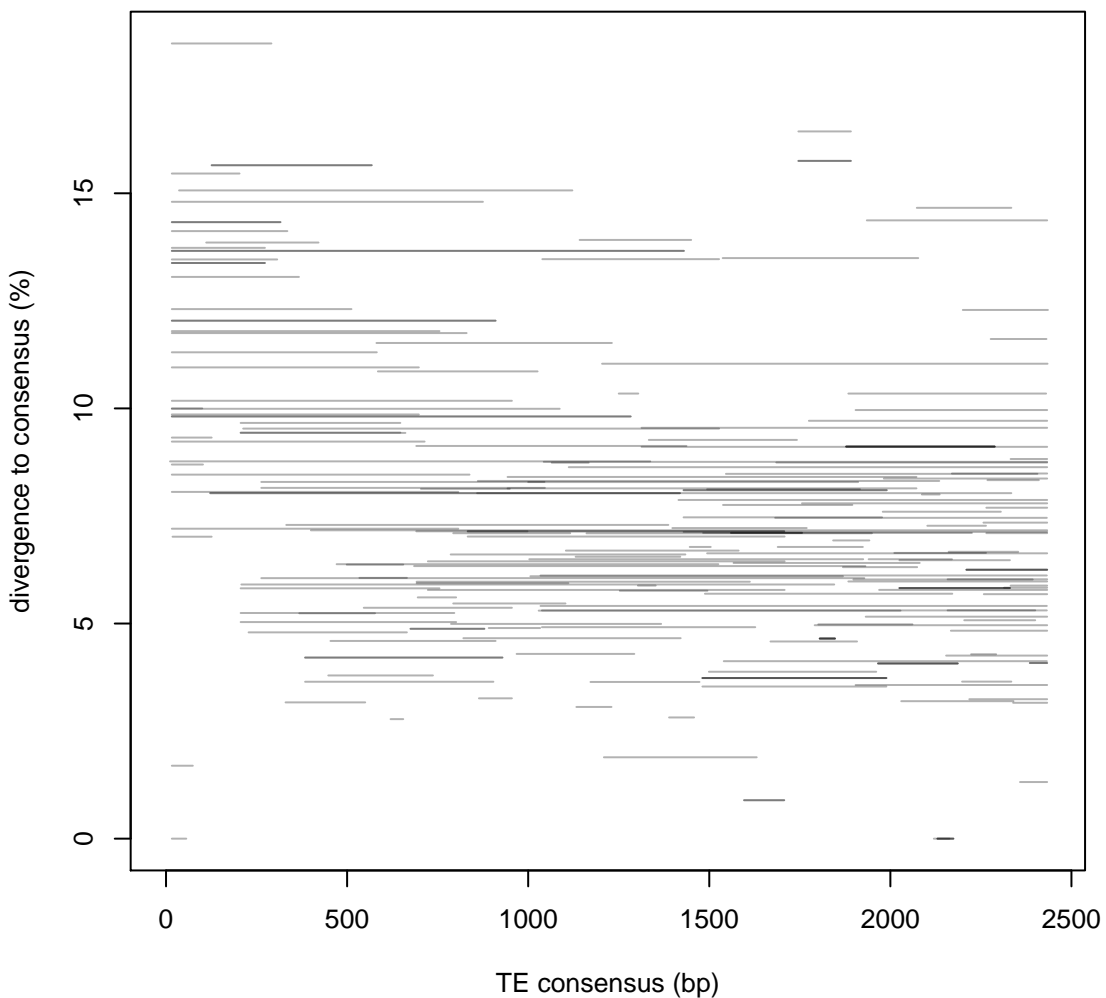
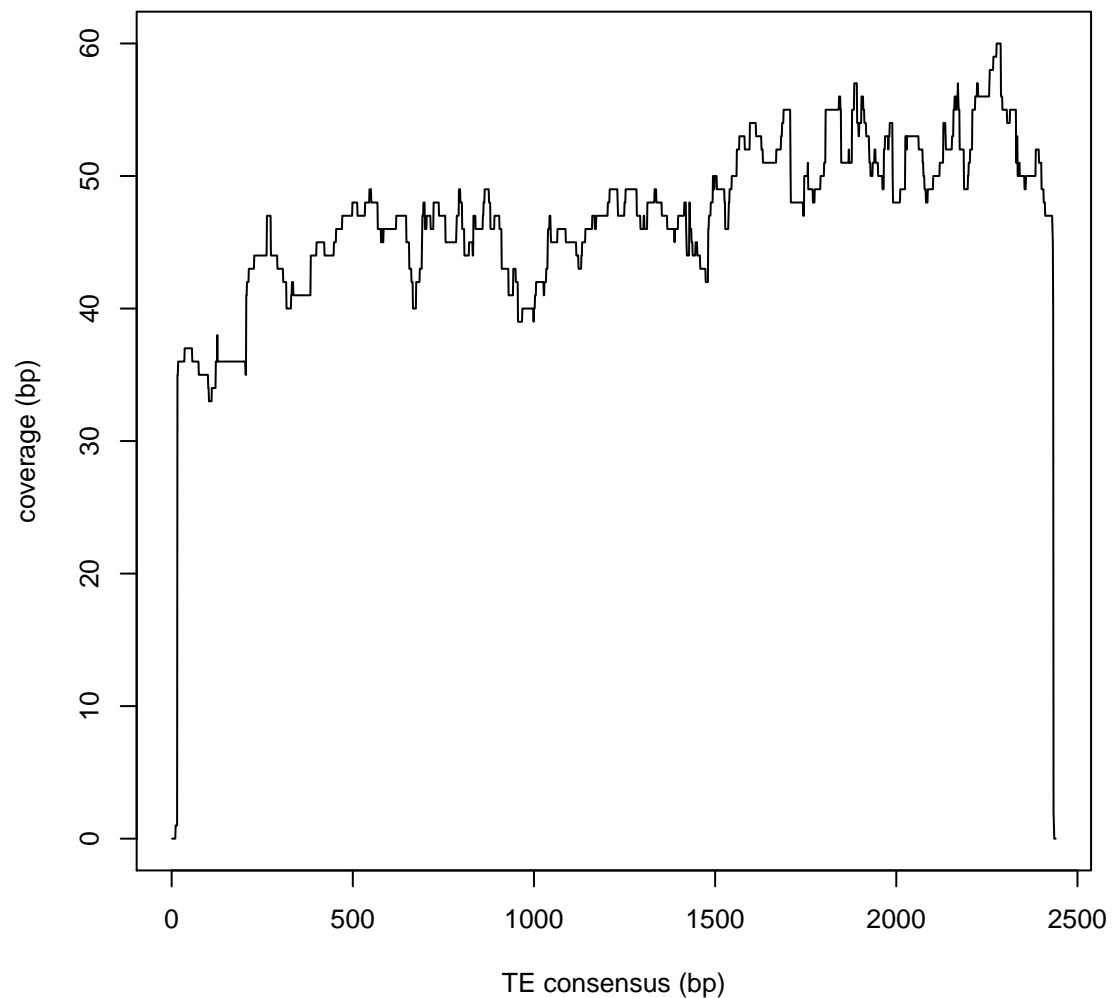


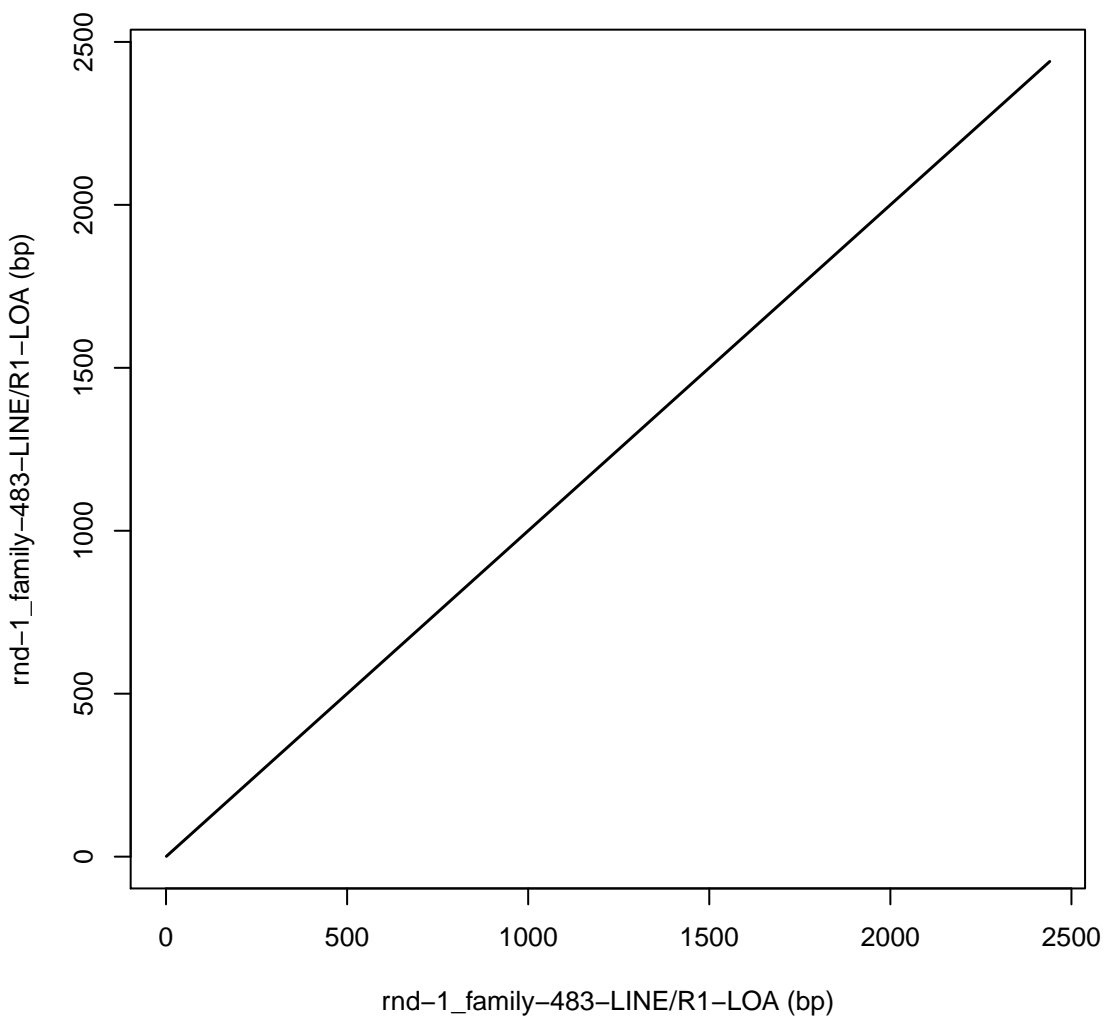
TE: rnd-1_family-483-LINE/R1-LOA
consensus size: 2440bp; fragments: 223; full length: 0 (>=2196bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

