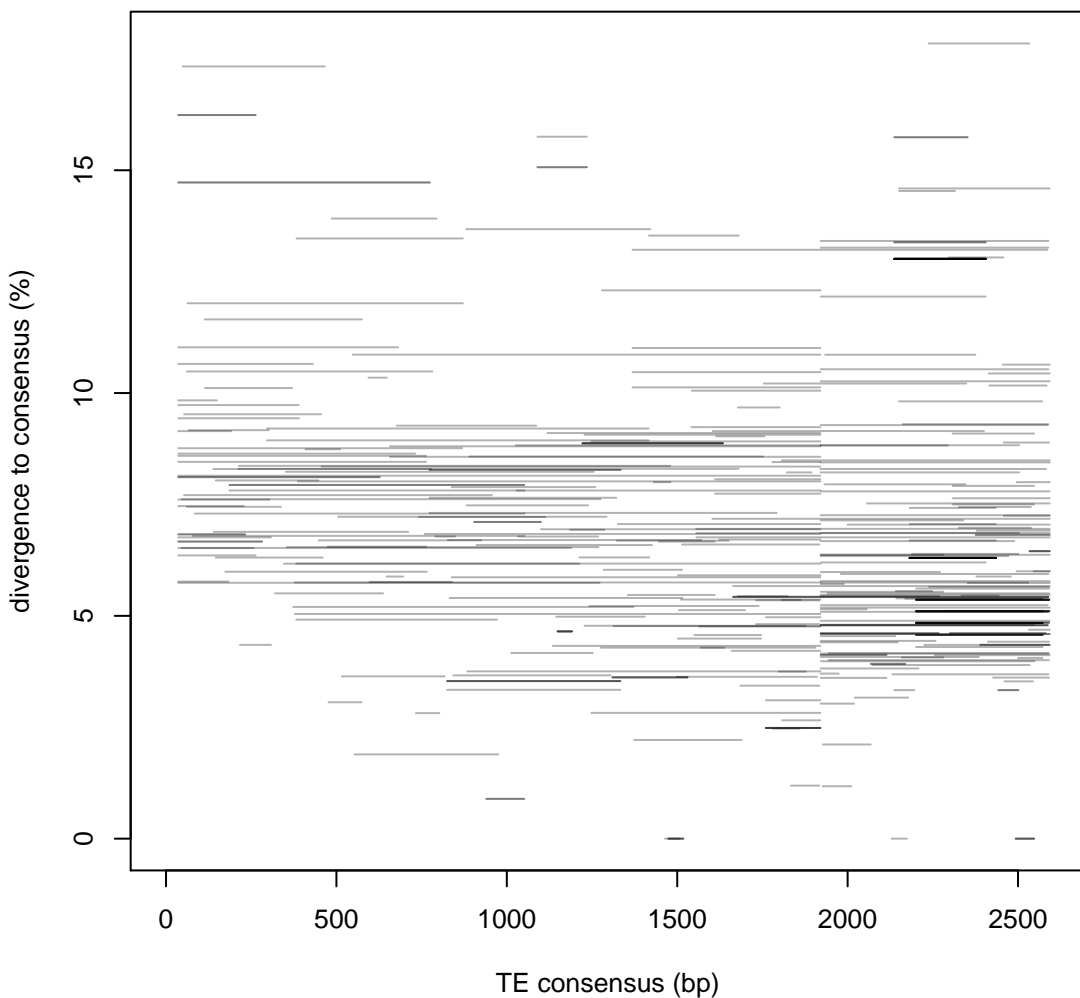
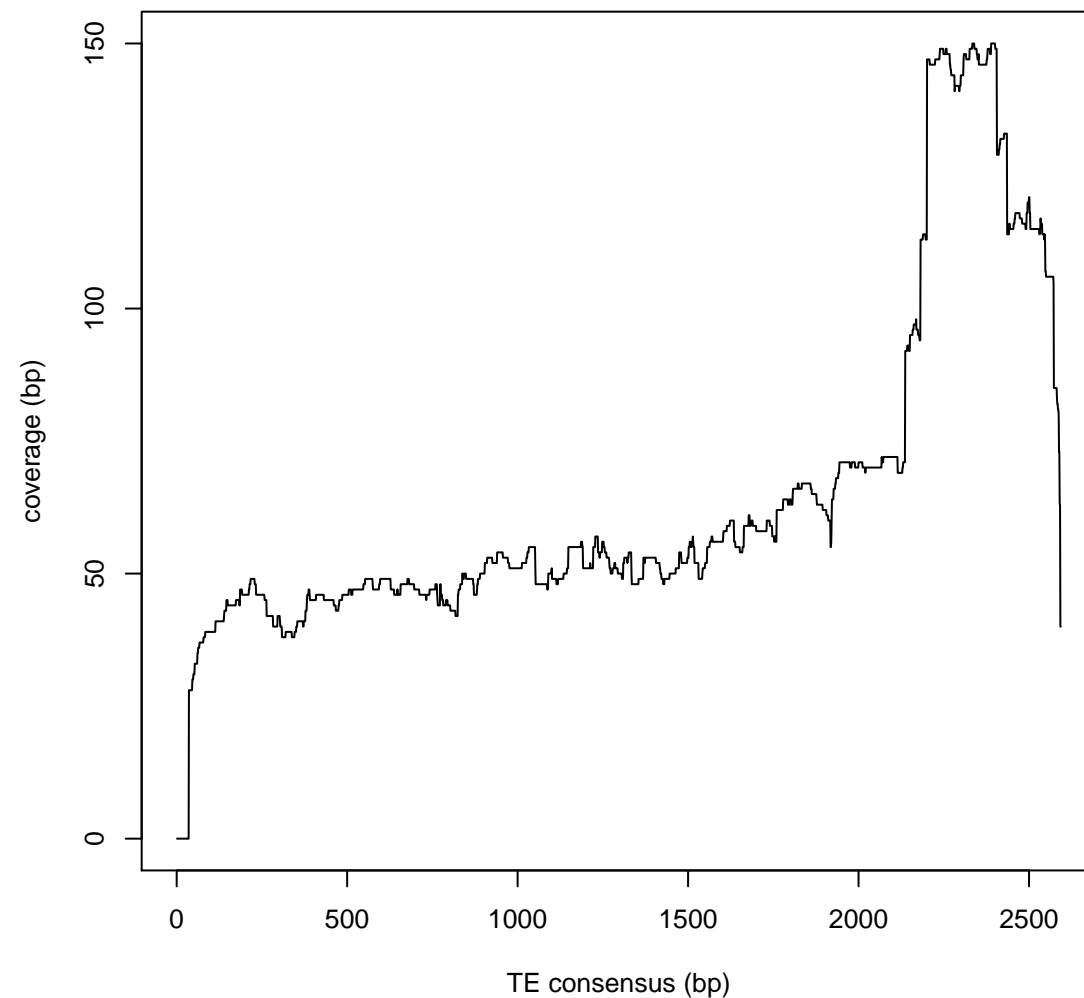


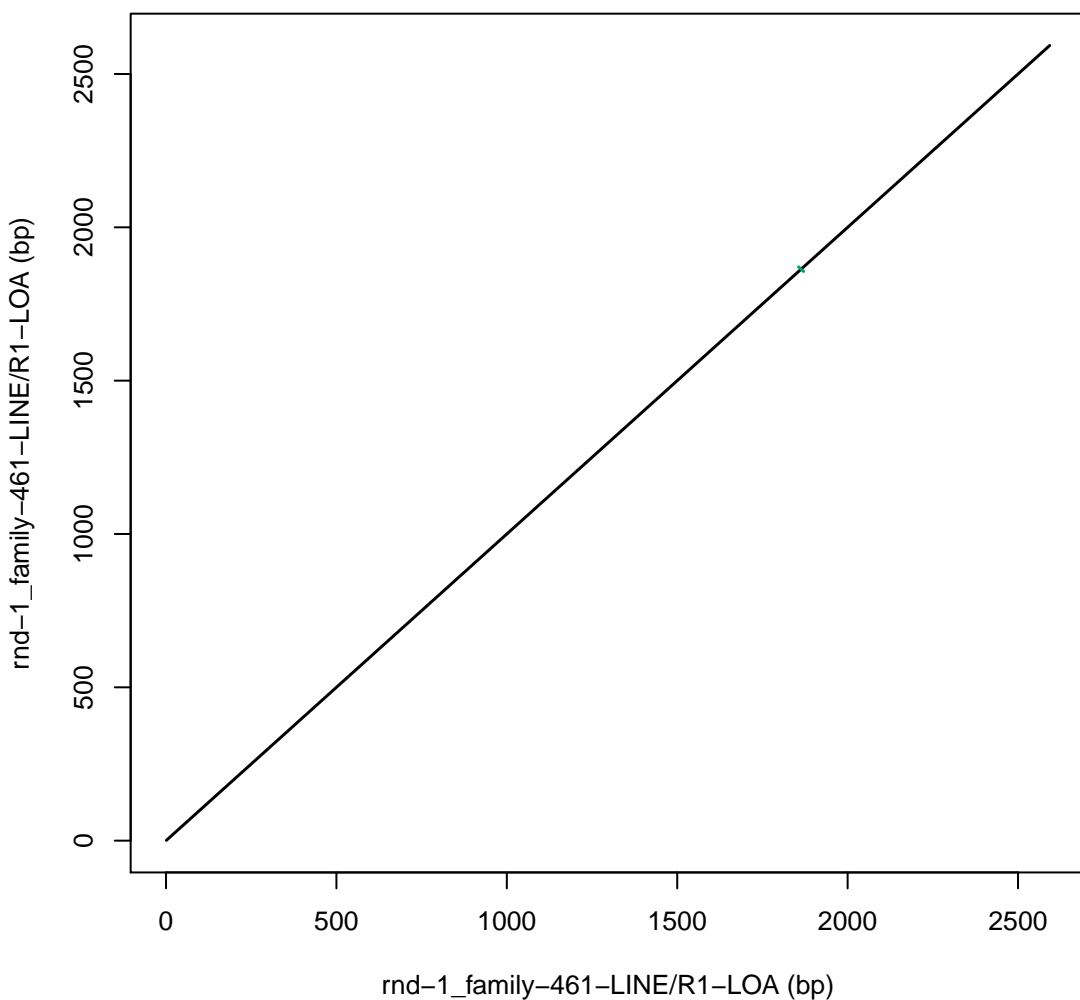
TE: rnd-1_family-461-LINE/R1-LOA
consensus size: 2593bp; fragments: 394; full length: 0 (>=2333.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

