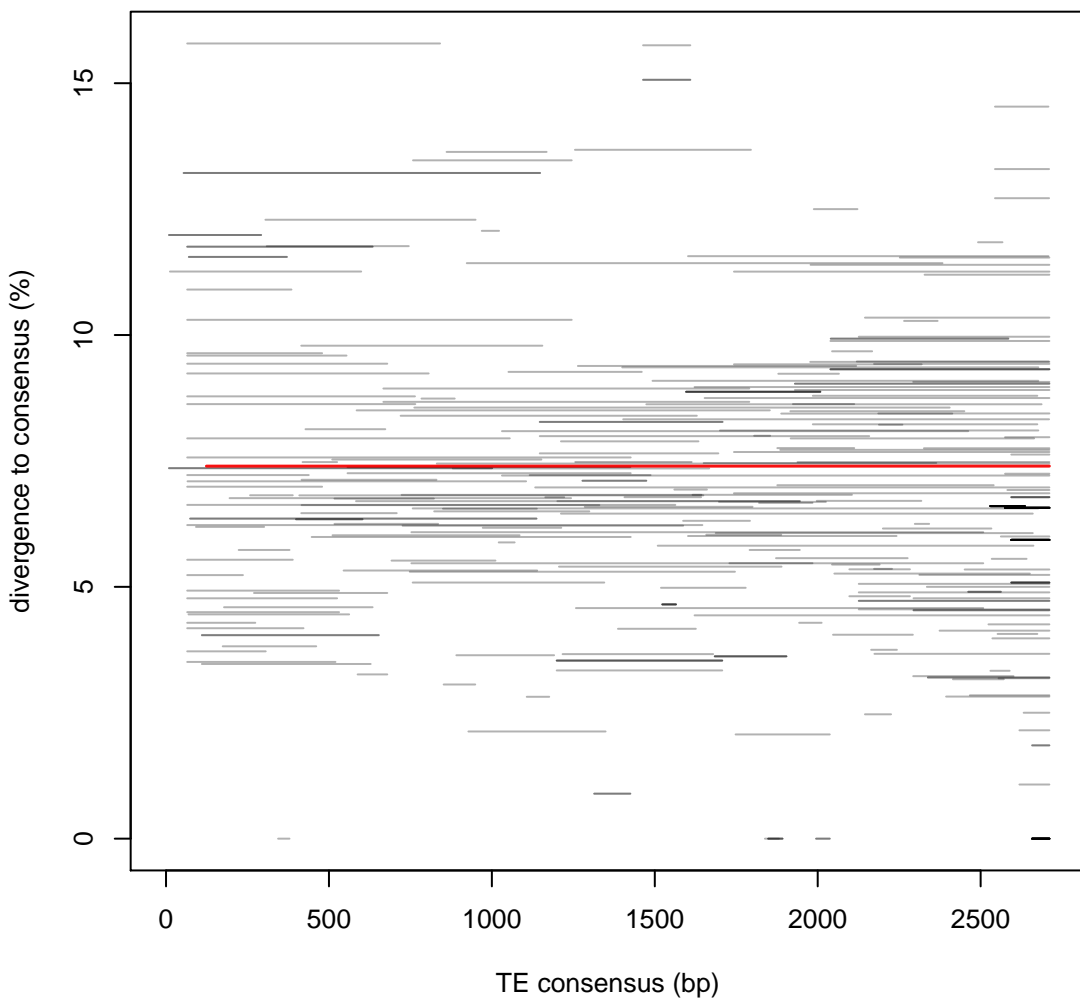
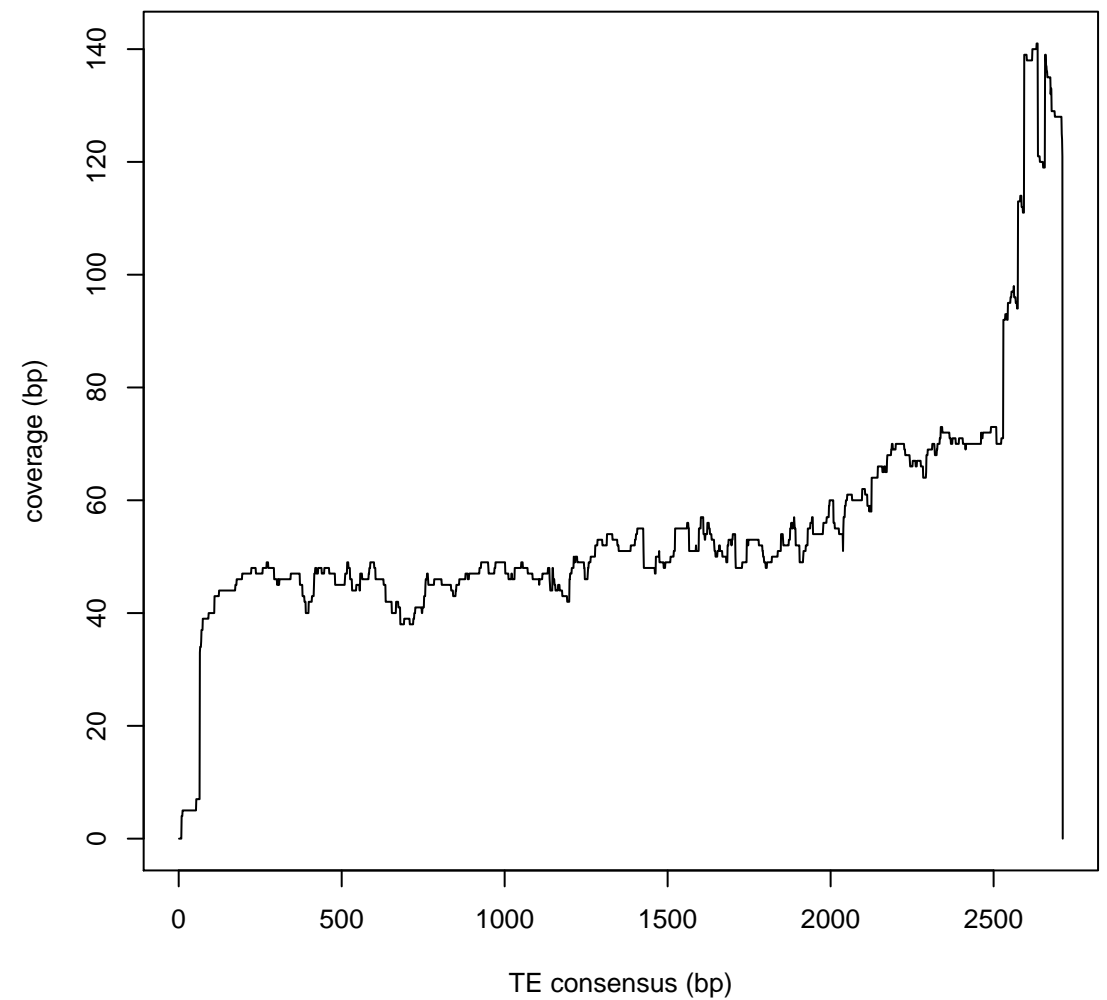


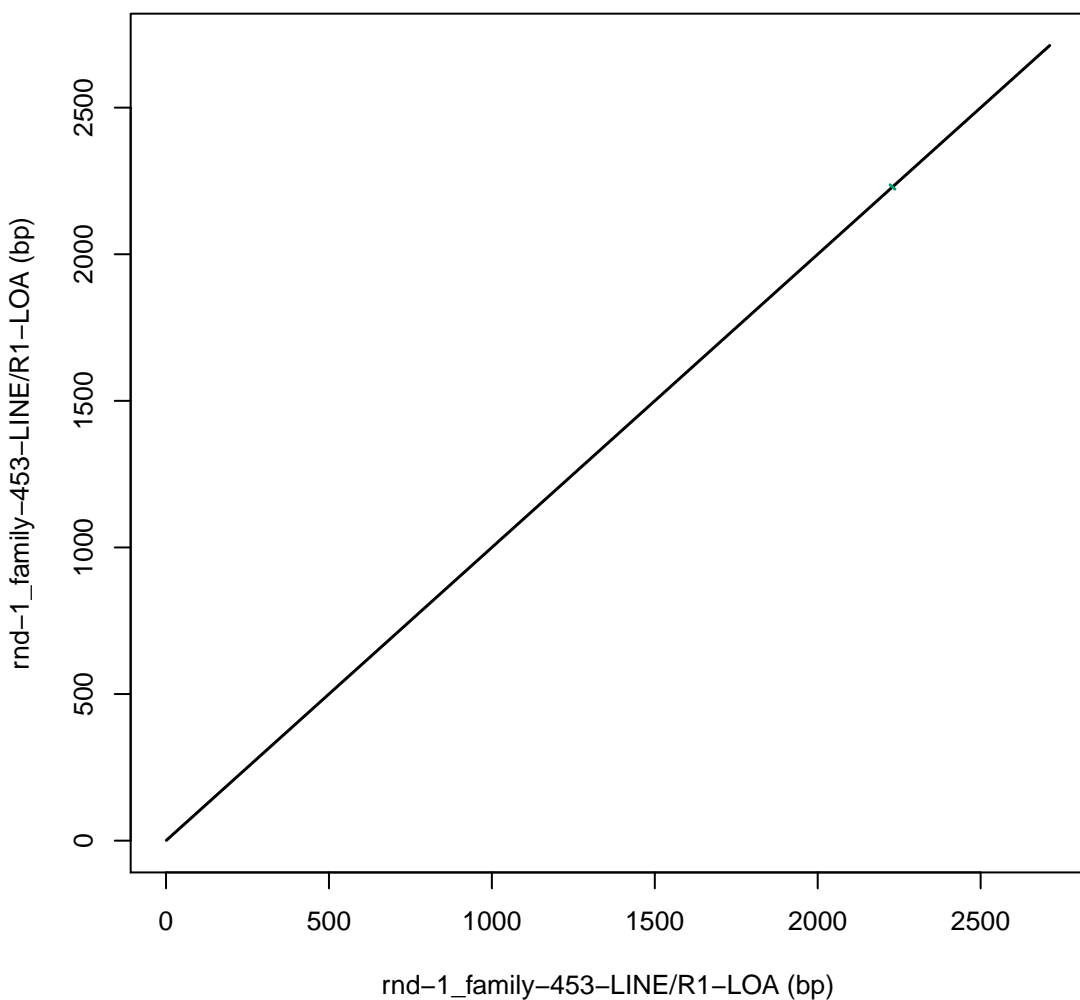
TE: rnd-1_family-453-LINE/R1-LOA
consensus size: 2712bp; fragments: 351; full length: 1 (≥ 2440.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

