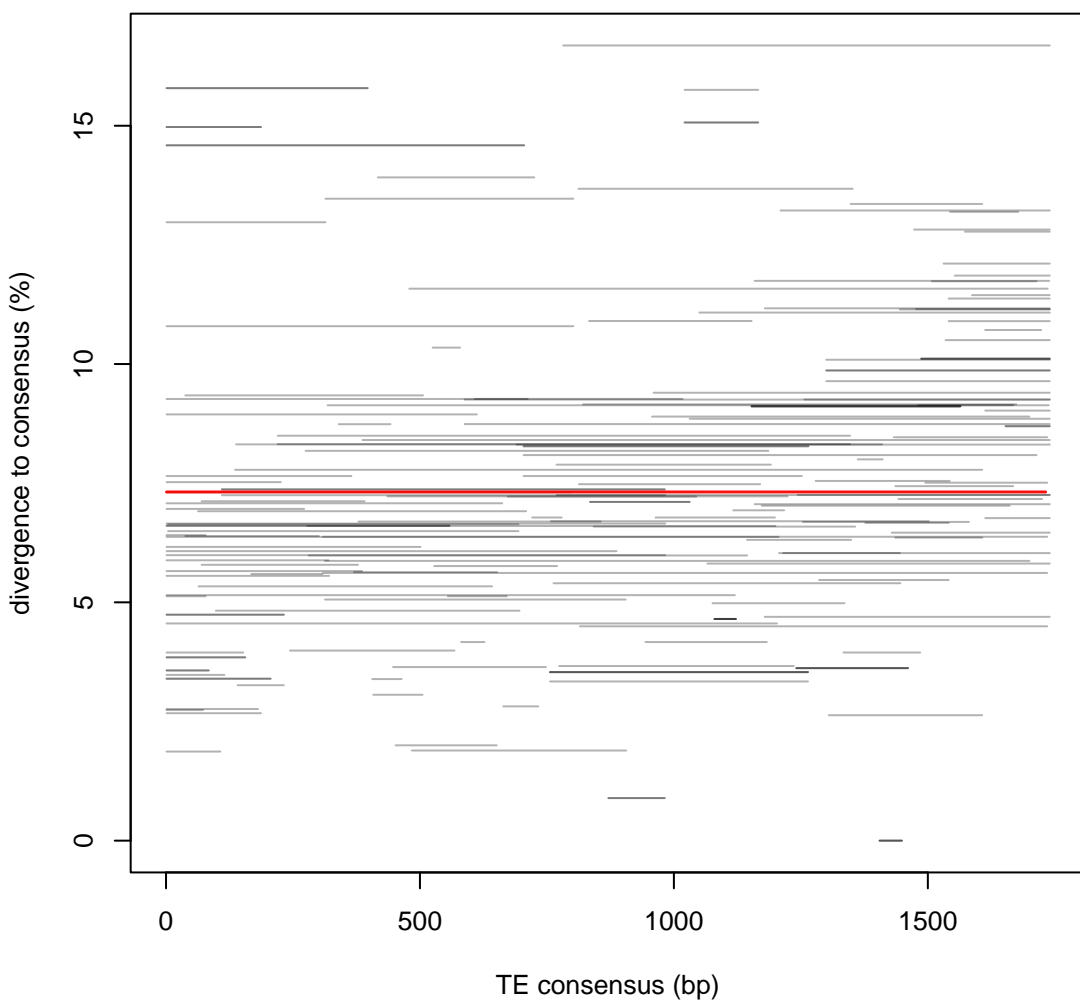
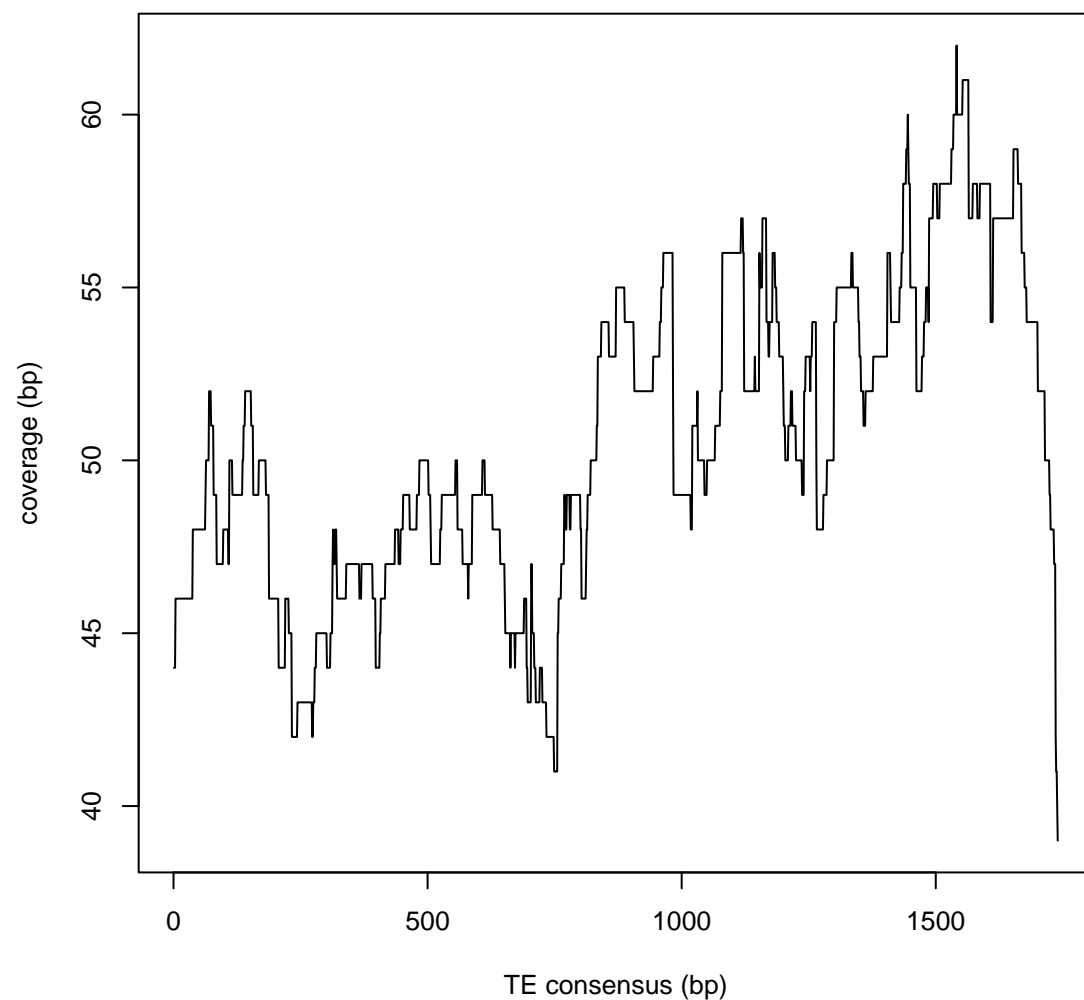


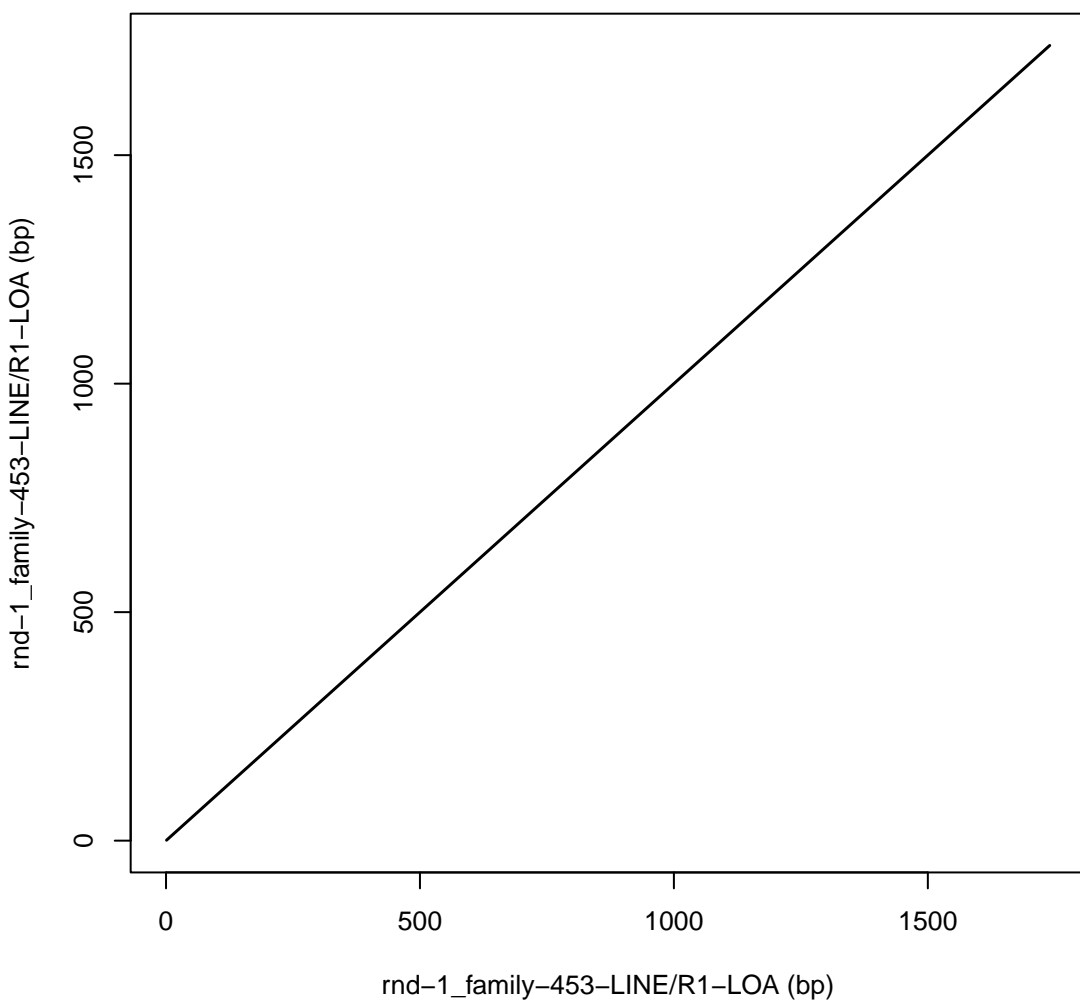
TE: rnd-1_family-453-LINE/R1-LOA
consensus size: 1740bp; fragments: 195; full length: 1 (>=1566bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

