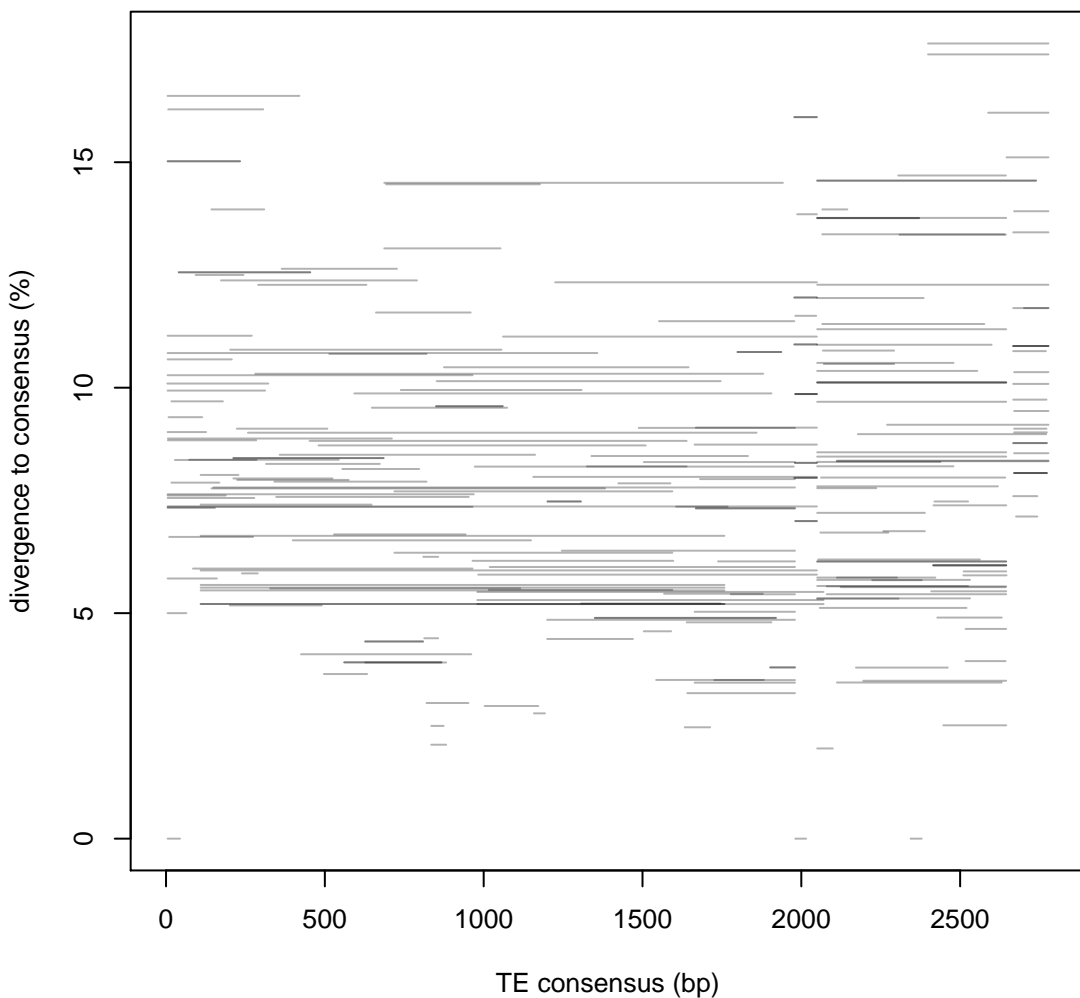
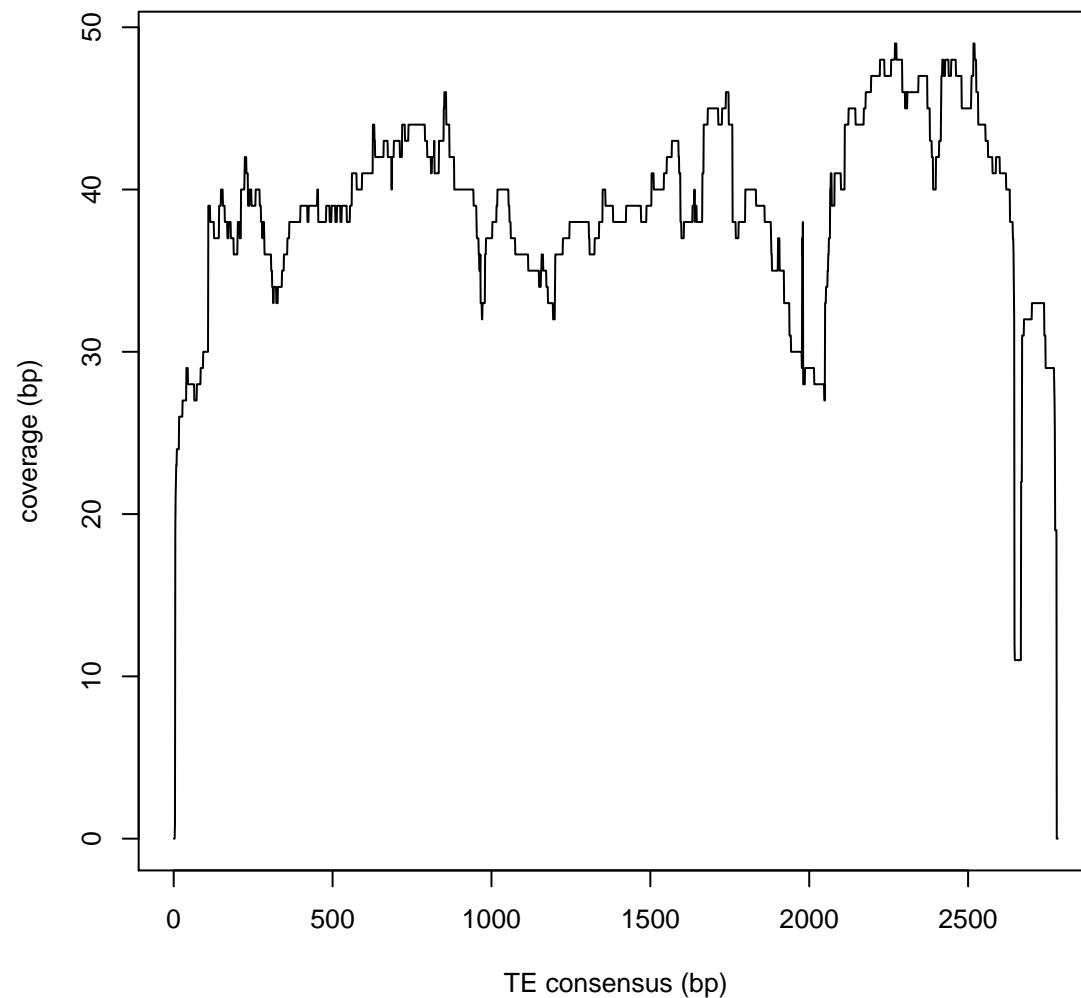


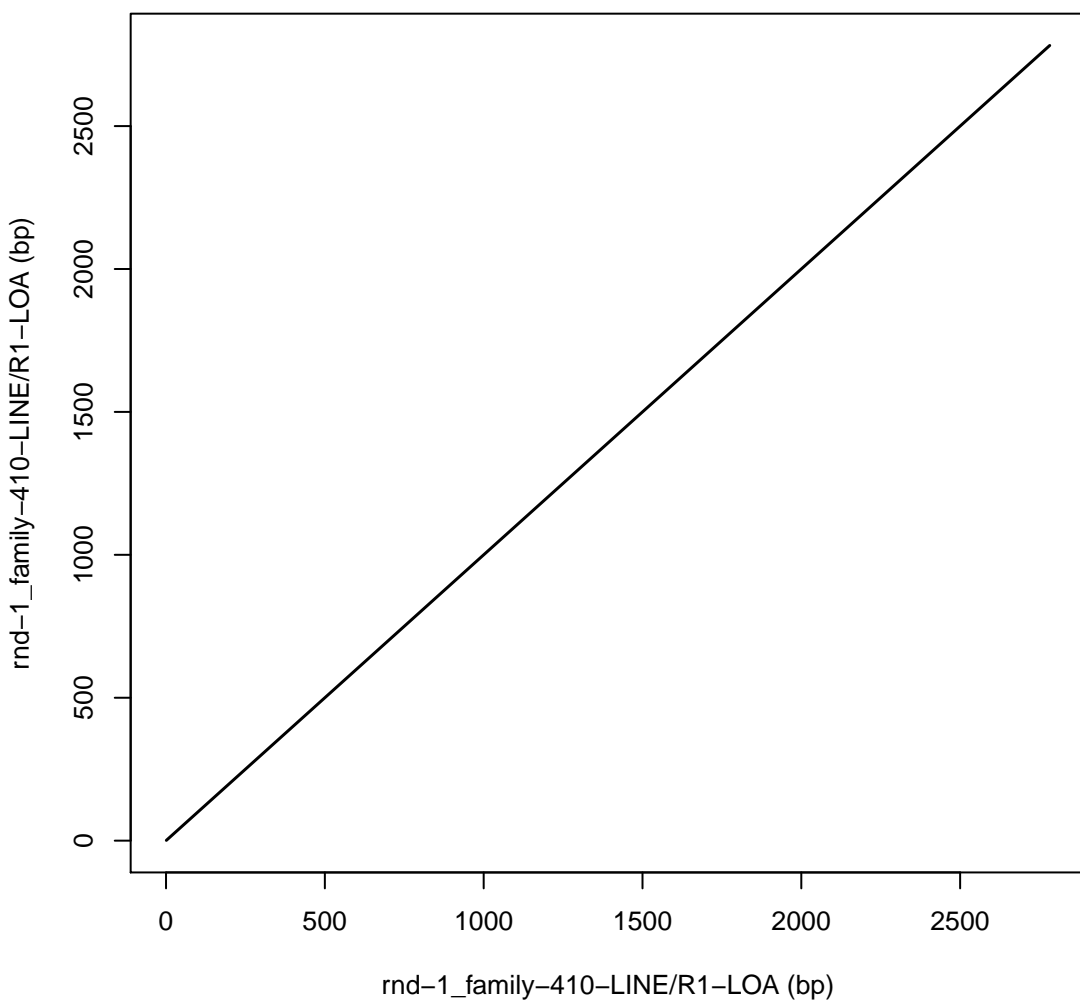
TE: rnd-1_family-410-LINE/R1-LOA
consensus size: 2782bp; fragments: 252; full length: 0 (>=2503.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

