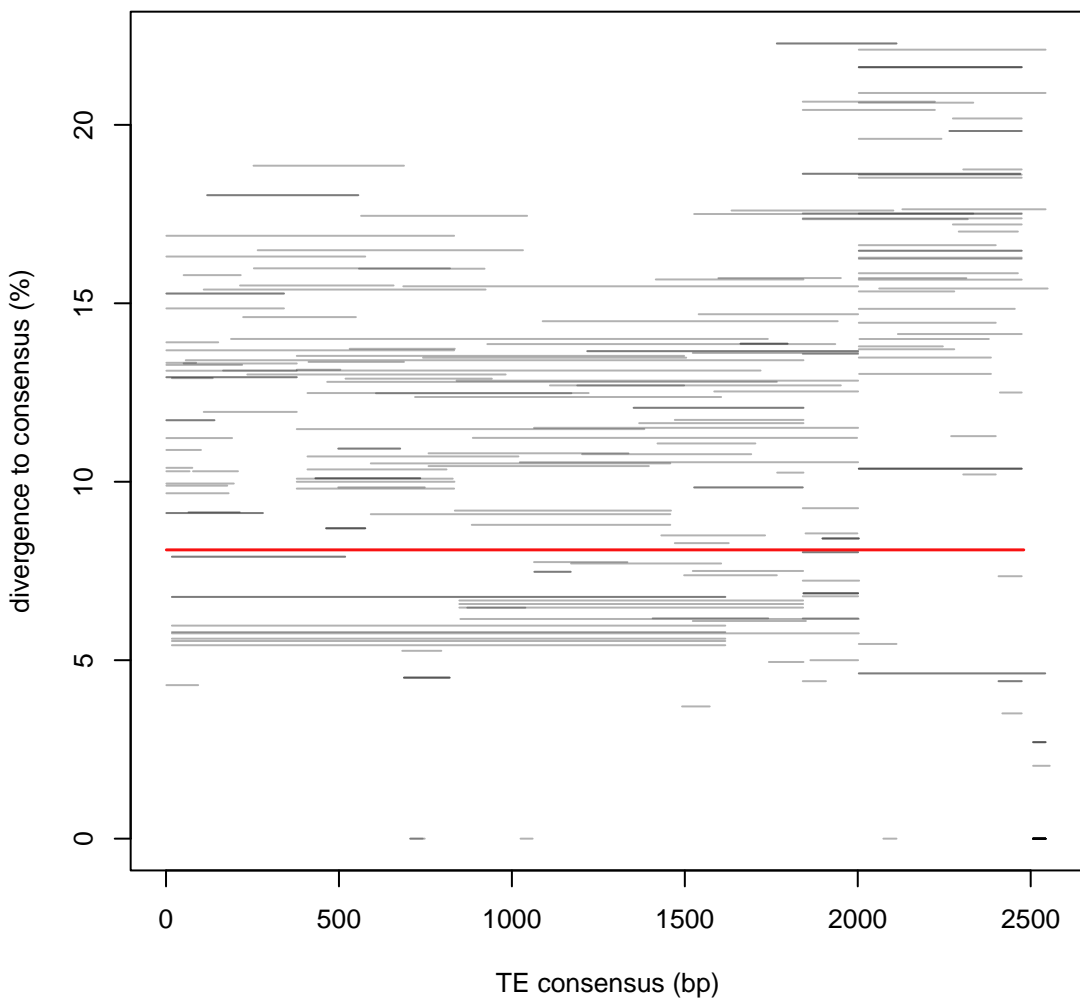
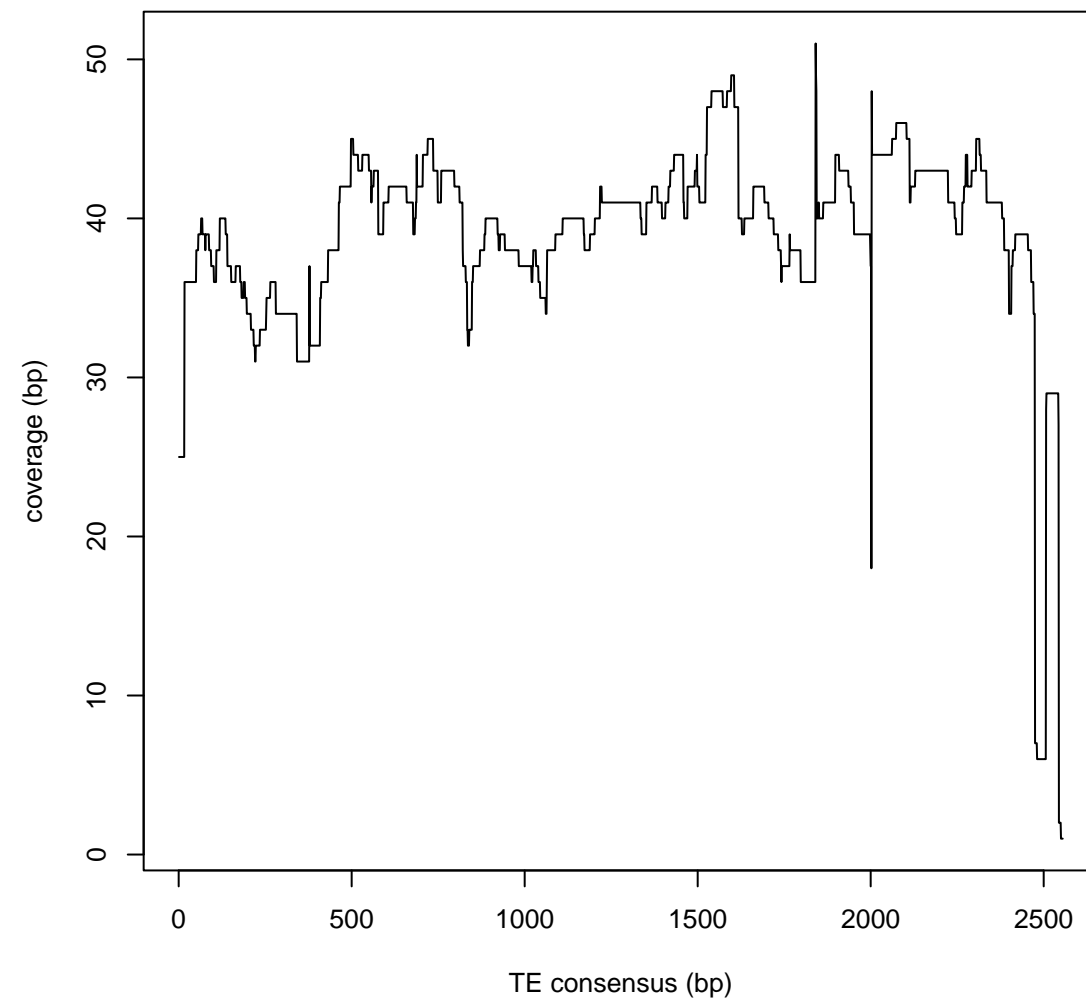


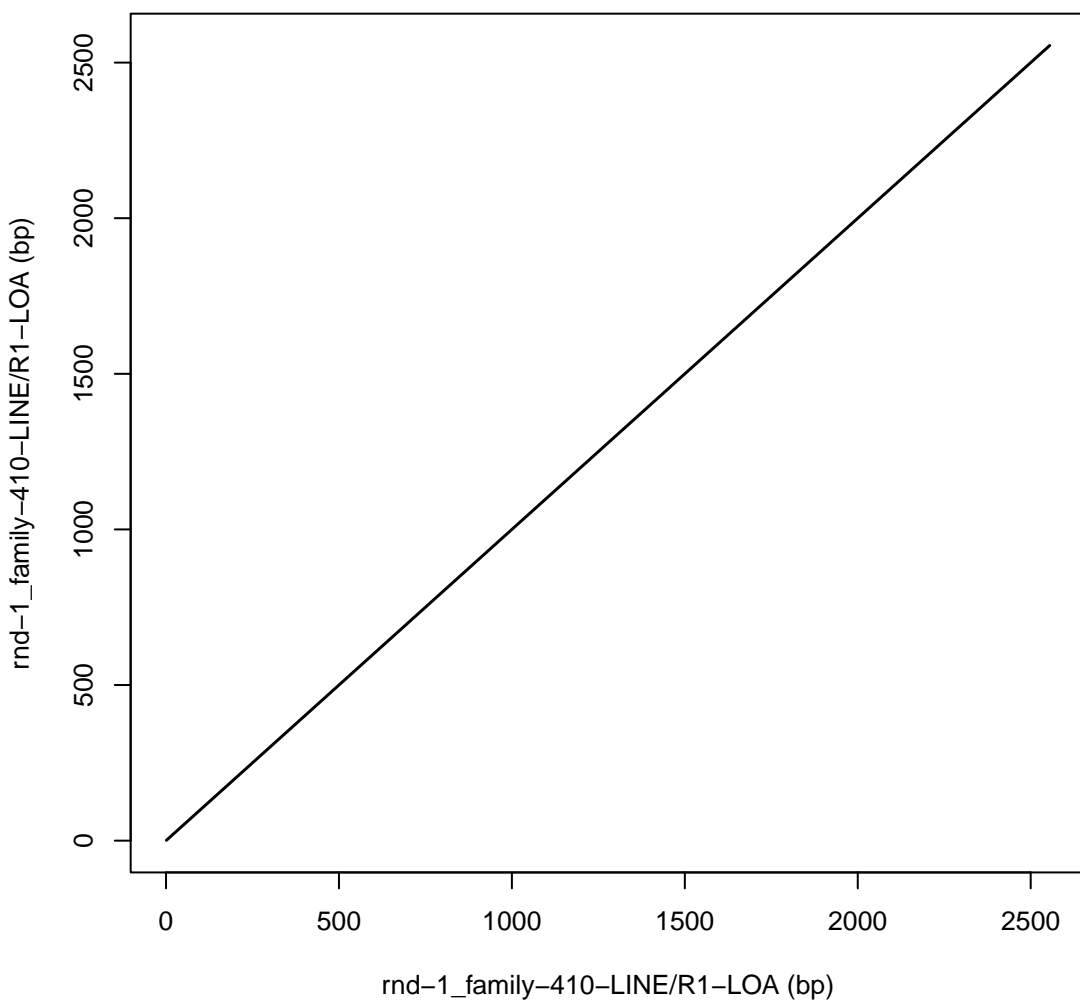
TE: rnd-1_family-410-LINE/R1-LOA
consensus size: 2555bp; fragments: 235; full length: 1 (>=2299.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

