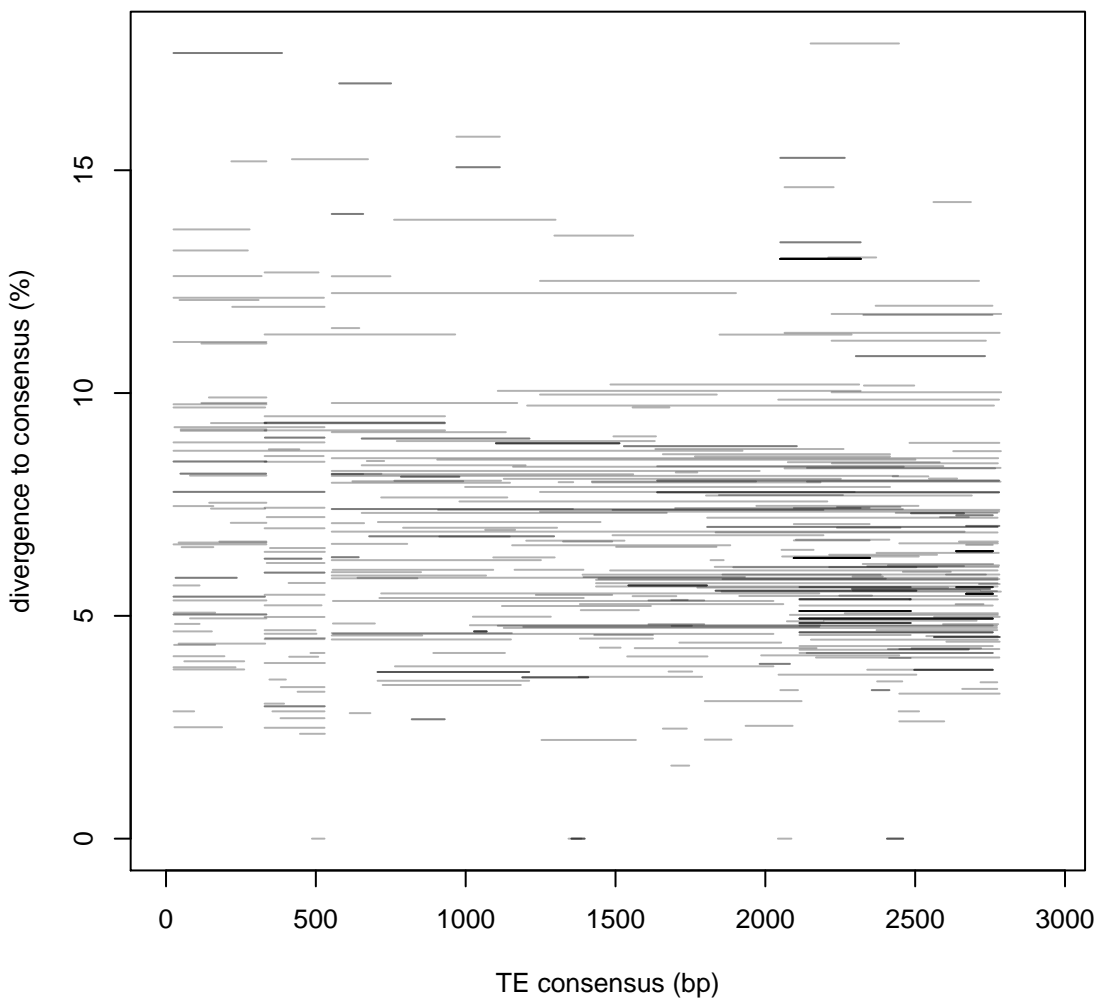
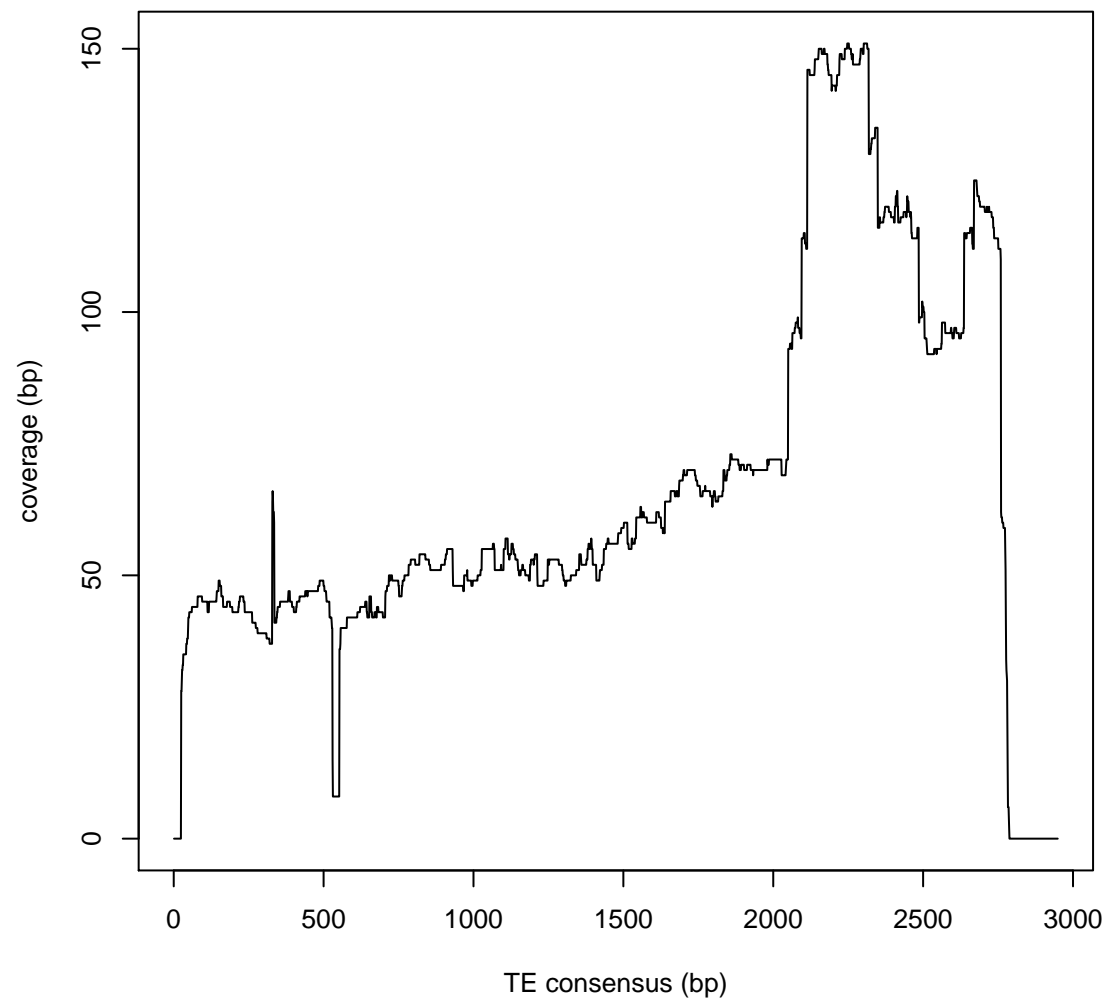


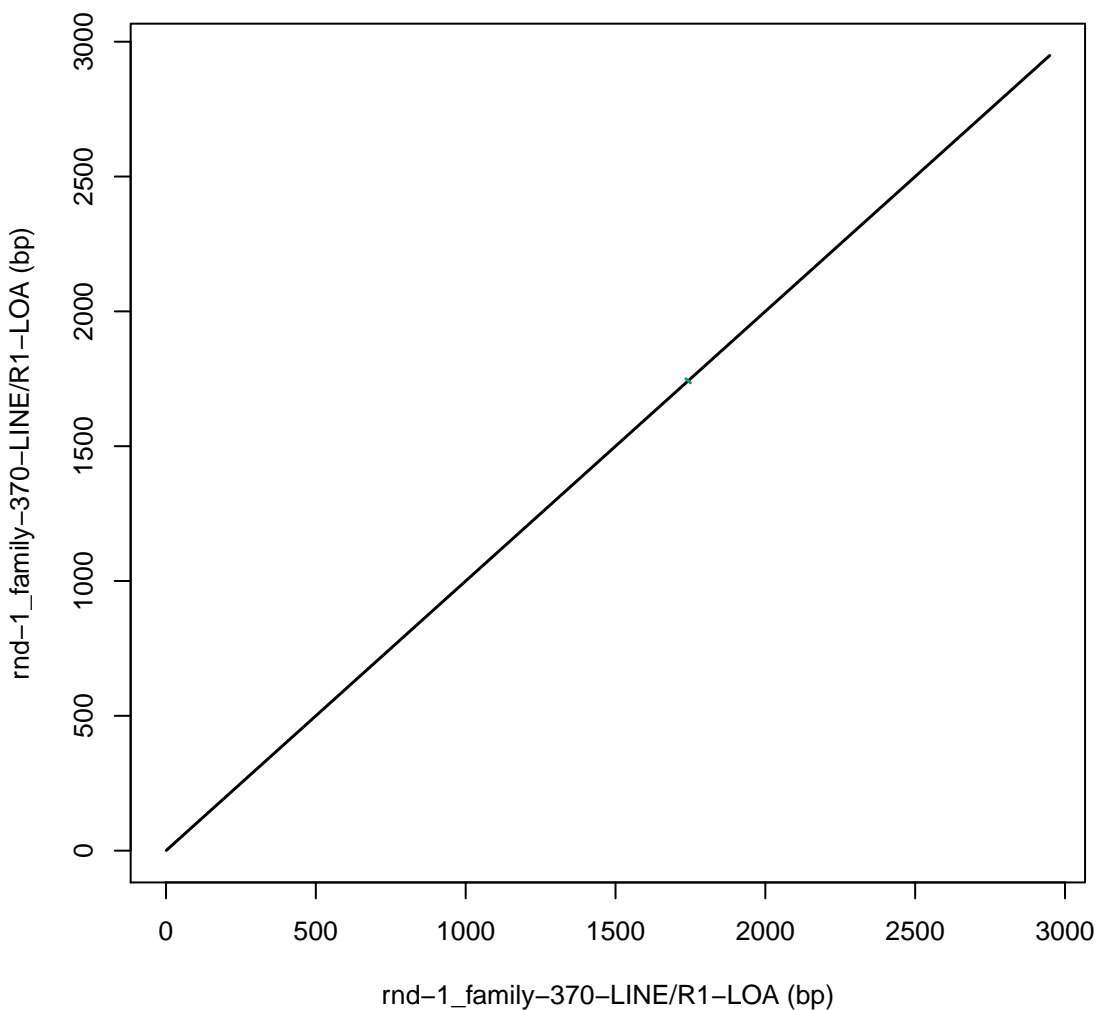
TE: rnd-1_family-370-LINE/R1-LOA
consensus size: 2949bp; fragments: 477; full length: 0 (>=2654.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

