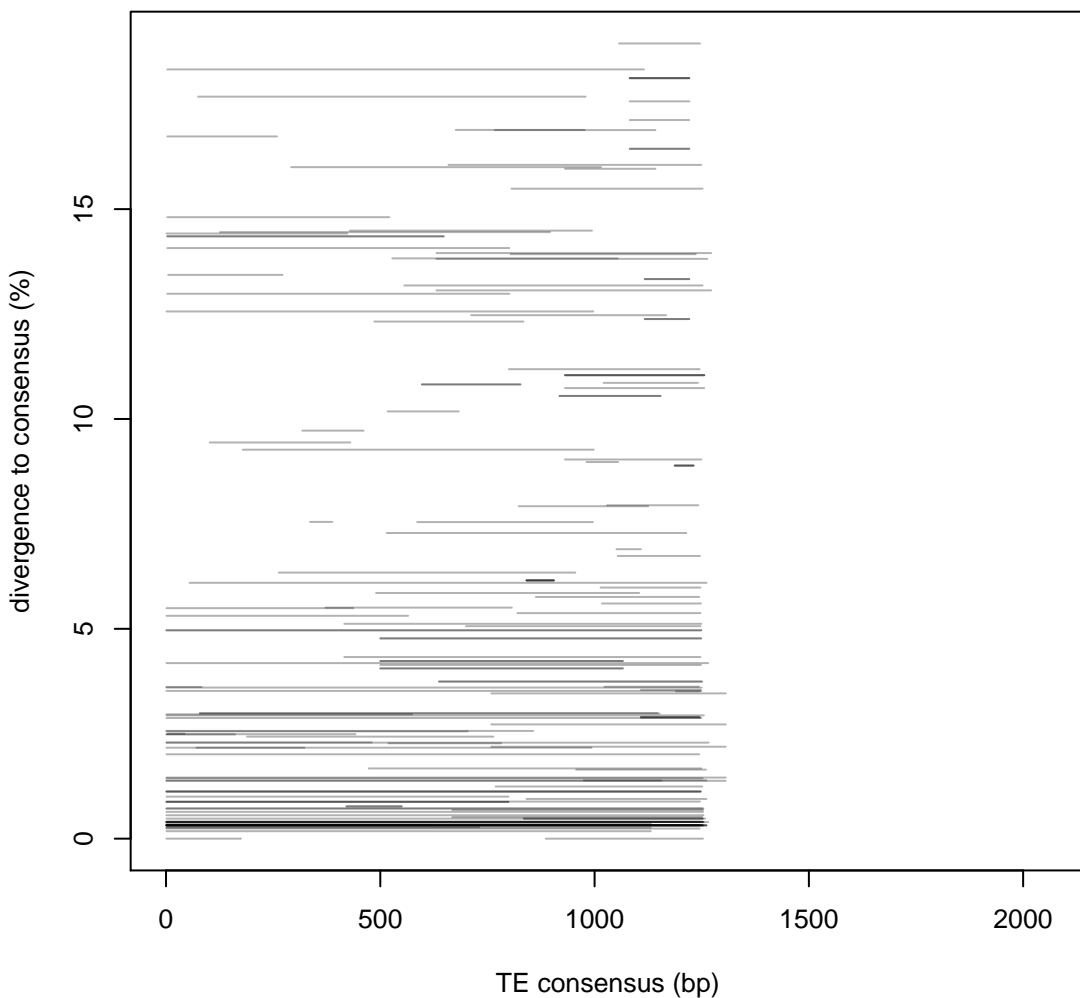
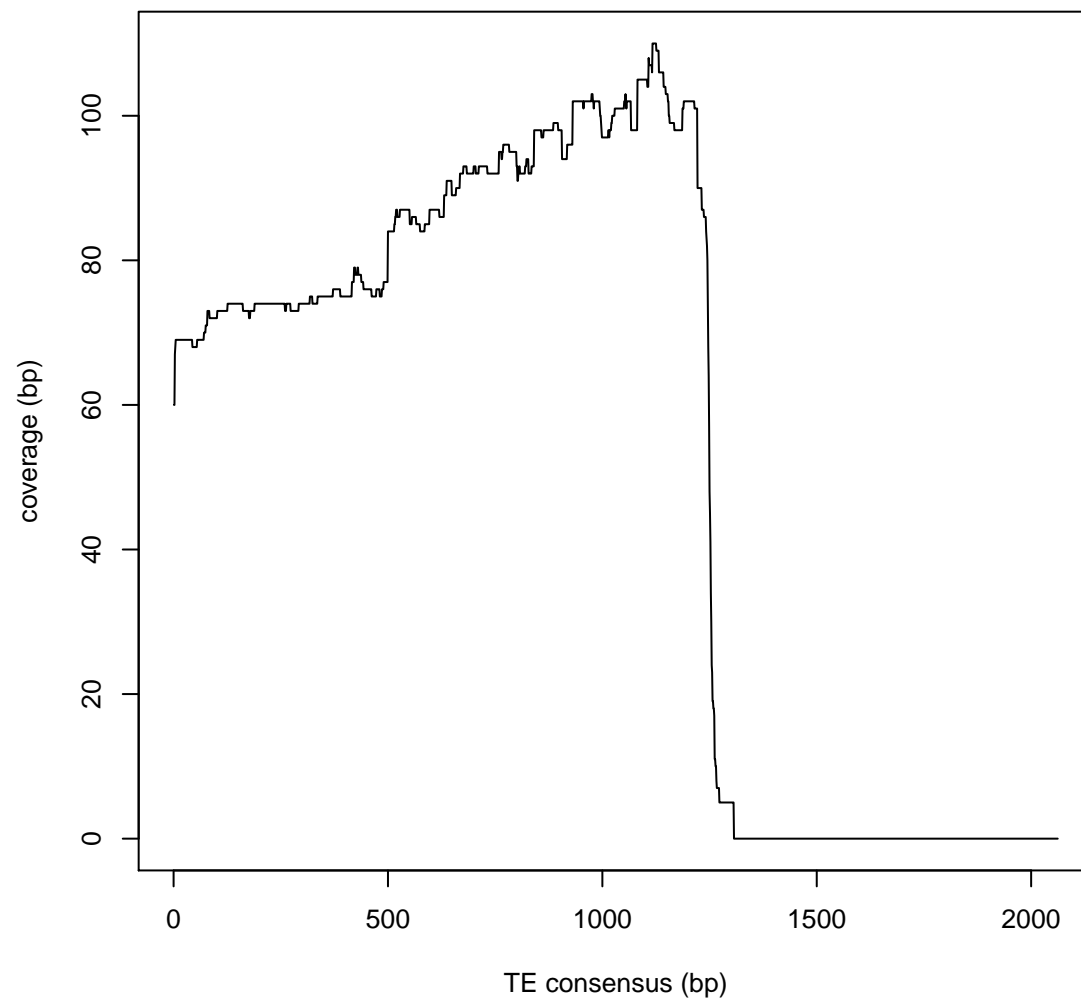


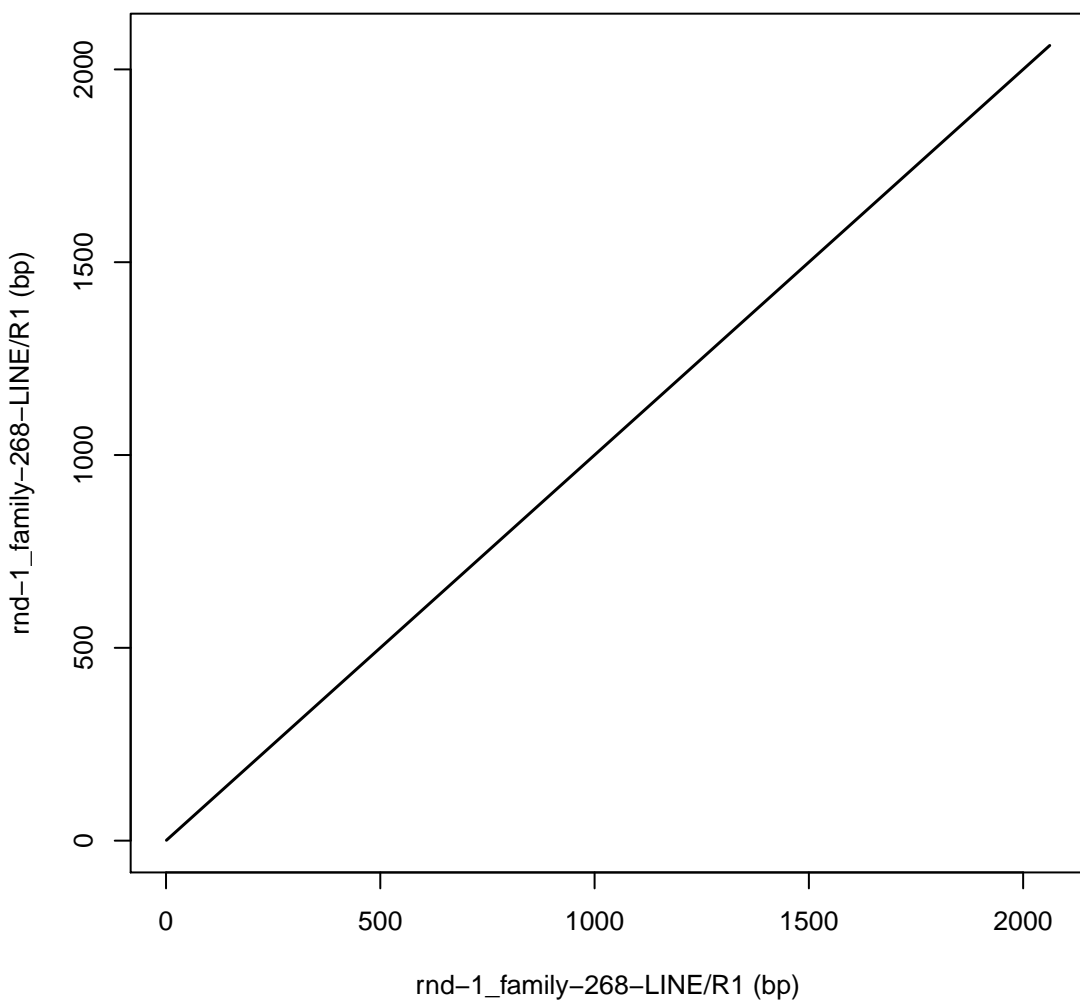
TE: rnd-1_family-268-LINE/R1
consensus size: 2062bp; fragments: 174; full length: 0 (≥ 1855.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

