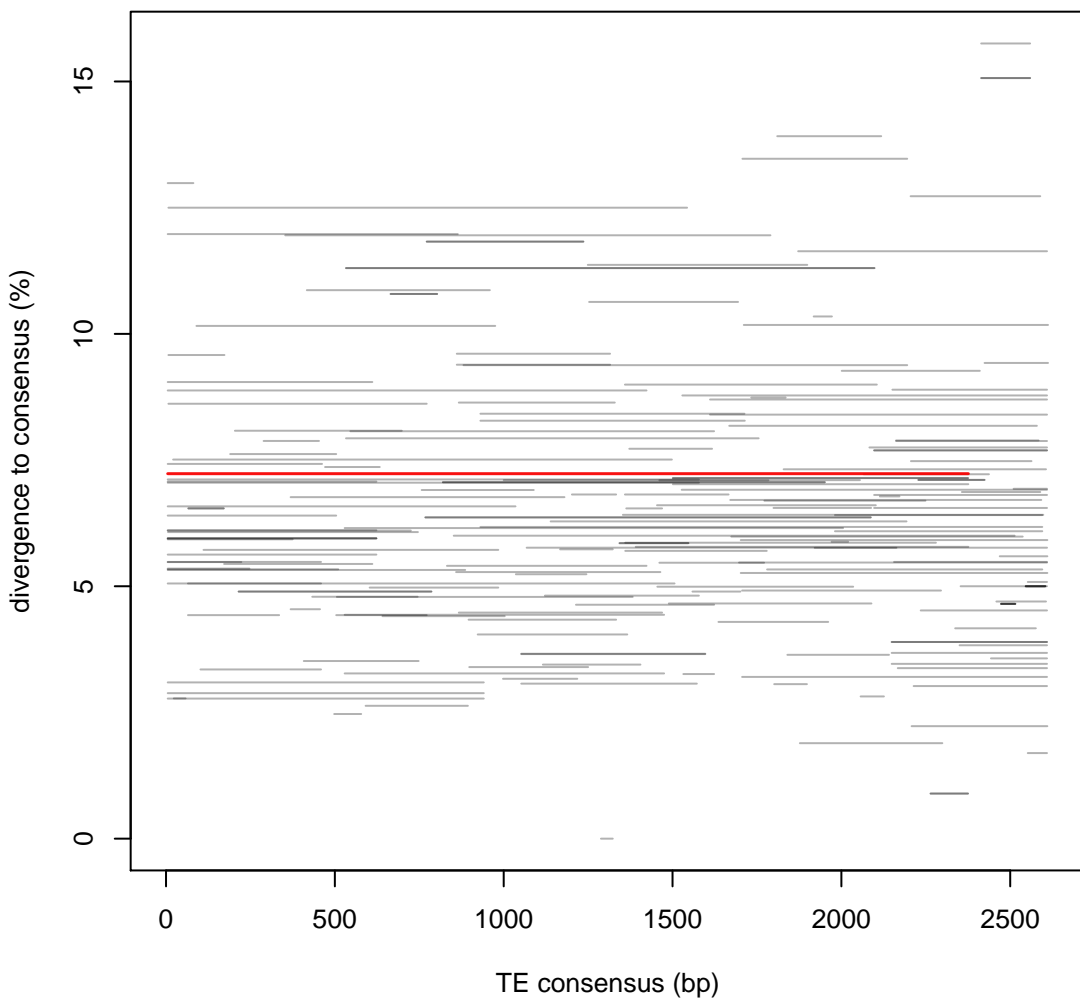
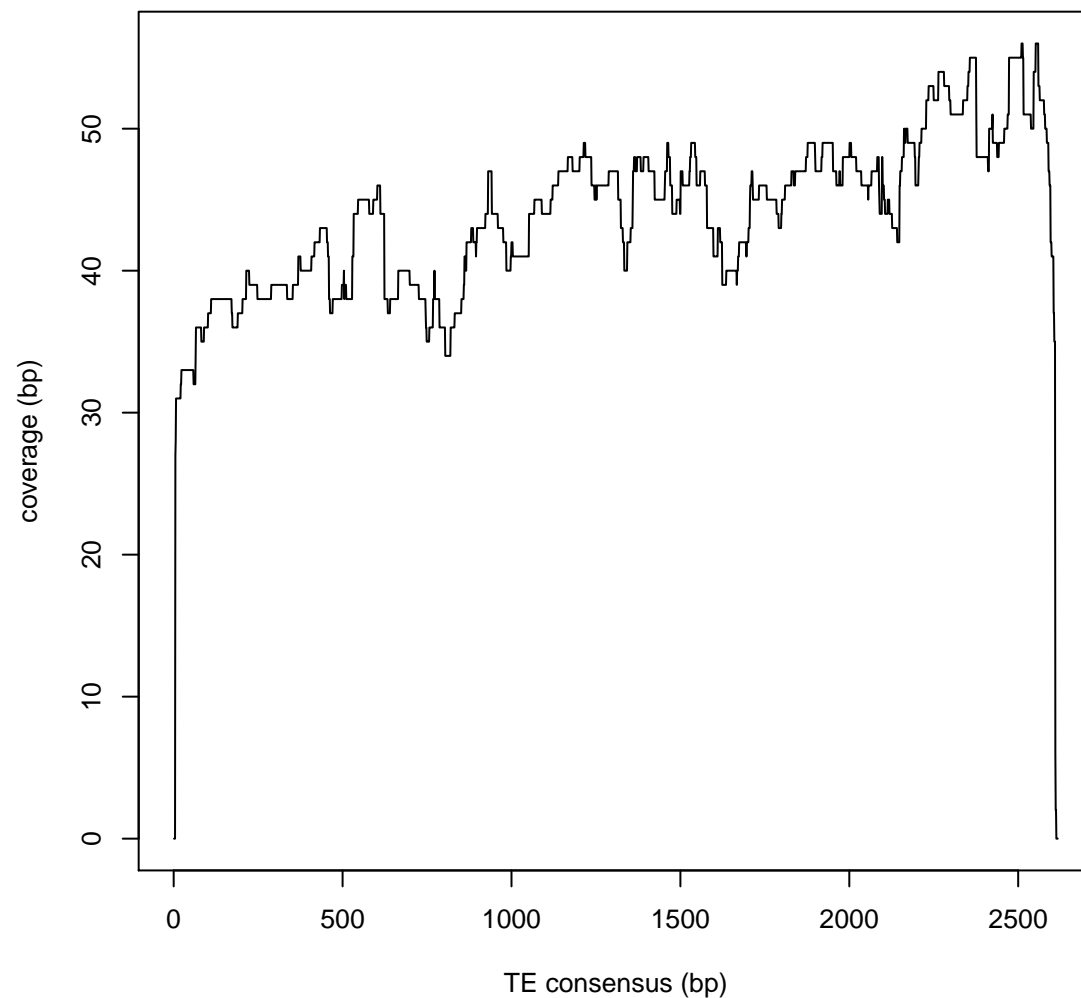


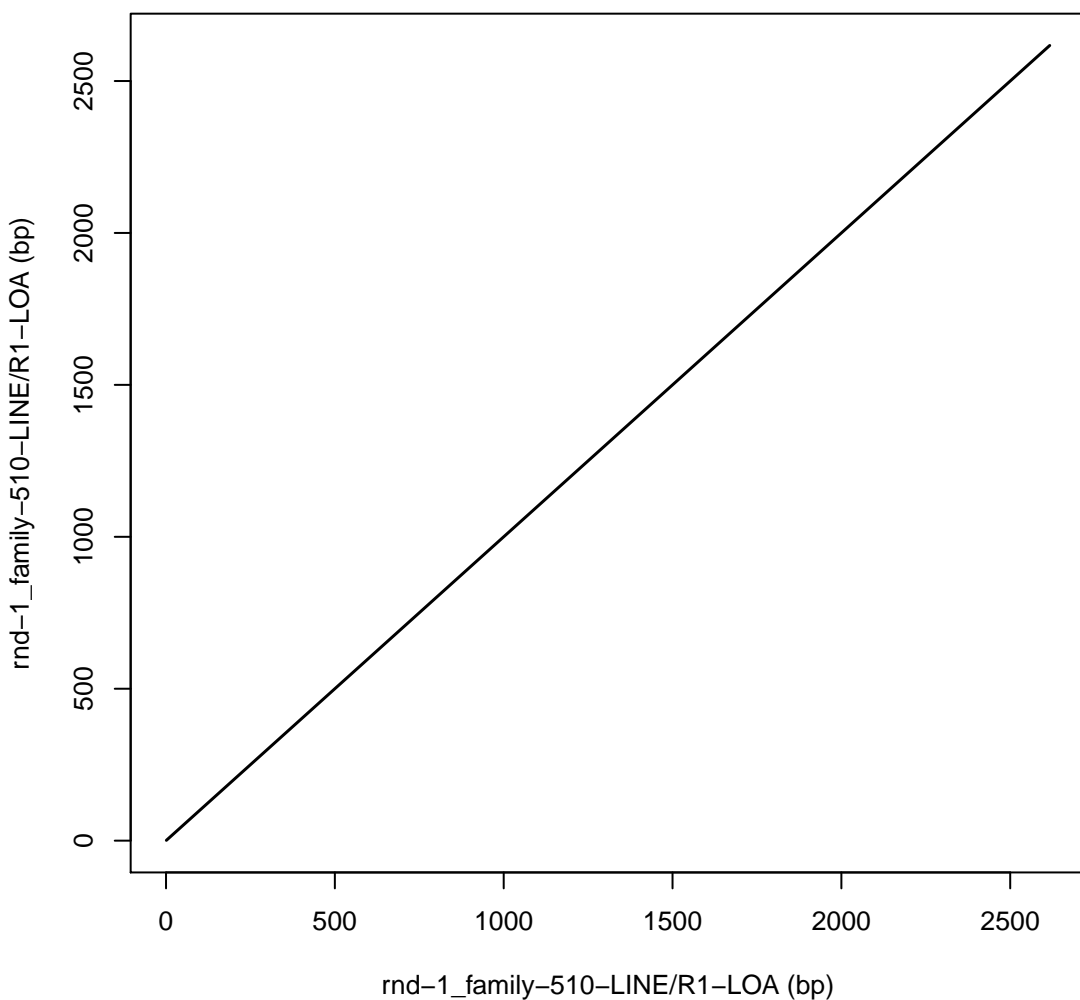
TE: rnd-1_family-510-LINE/R1-LOA
consensus size: 2617bp; fragments: 204; full length: 1 (>=2355.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

