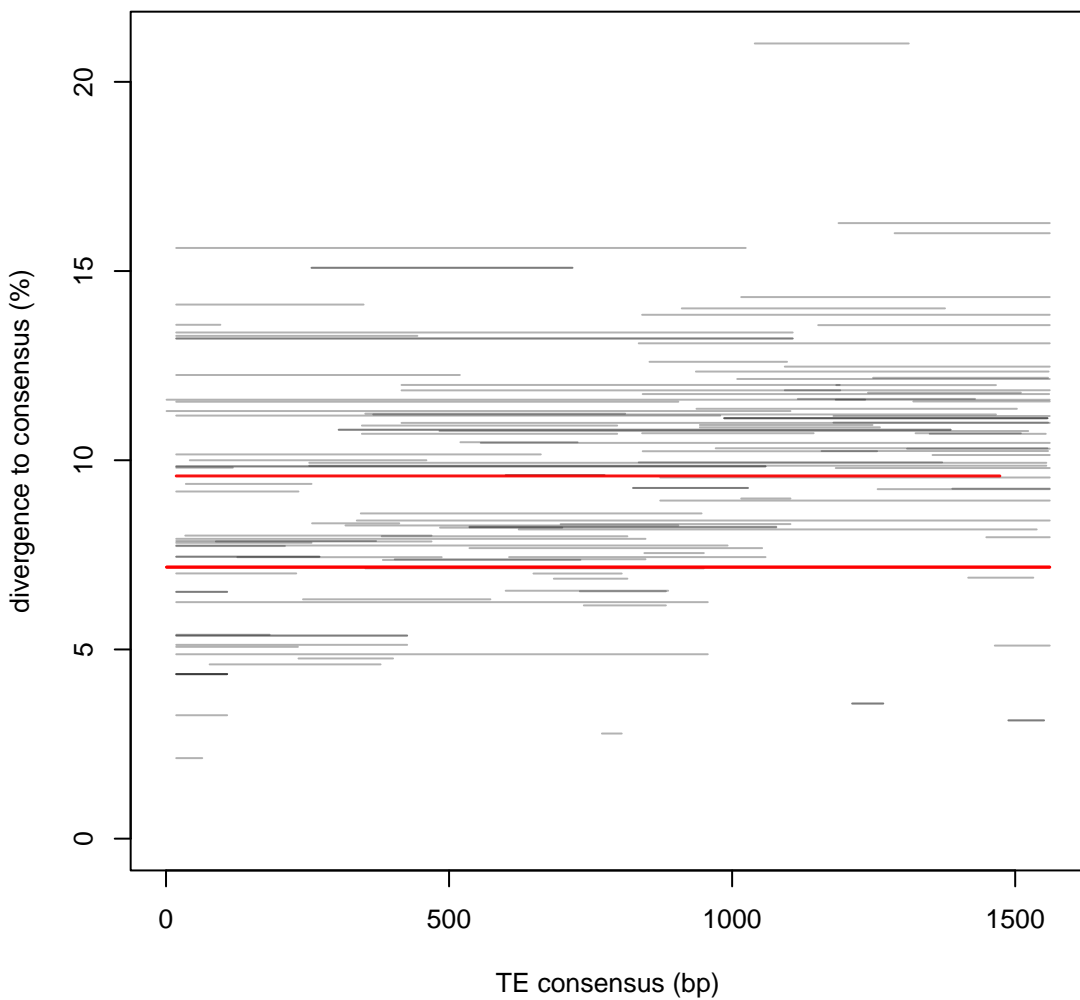
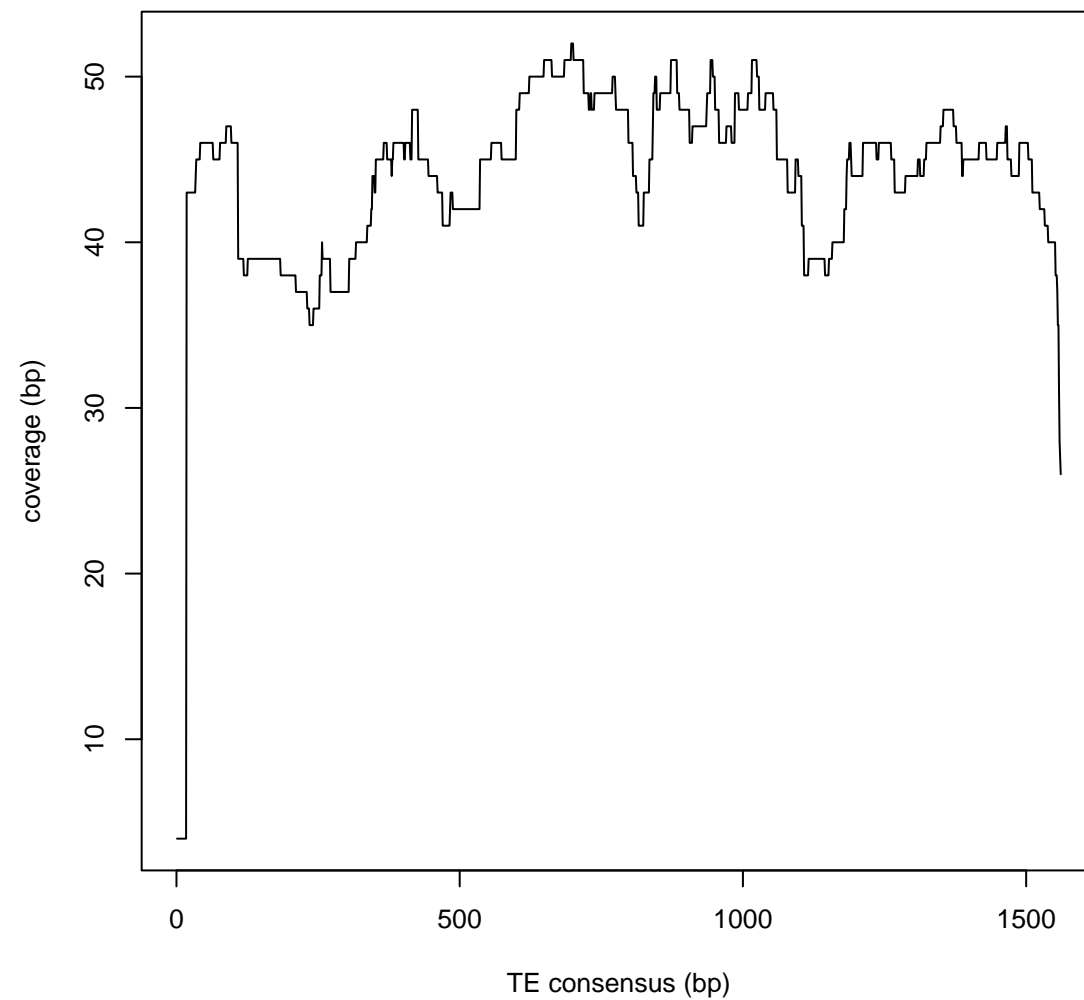


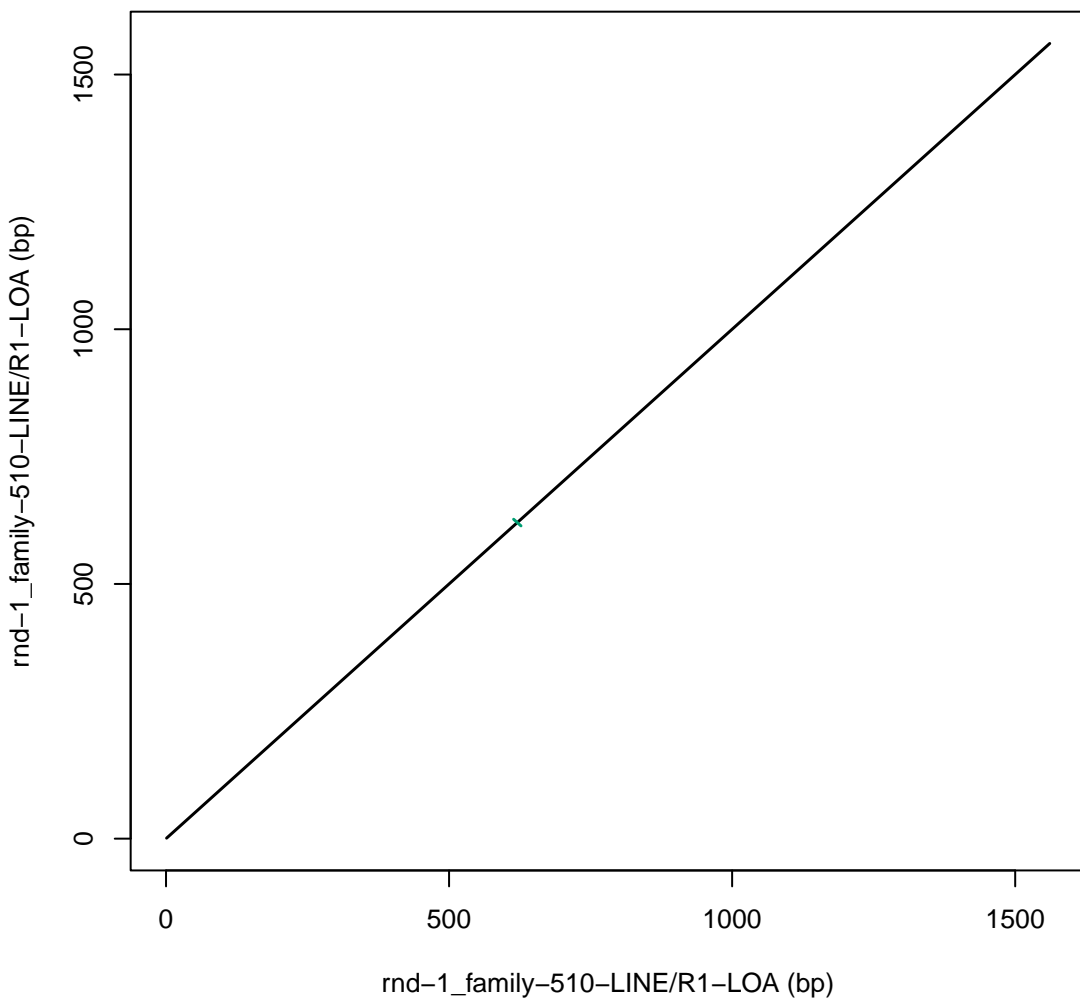
TE: rnd-1_family-510-LINE/R1-LOA
consensus size: 1561bp; fragments: 143; full length: 3 (>=1404.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

