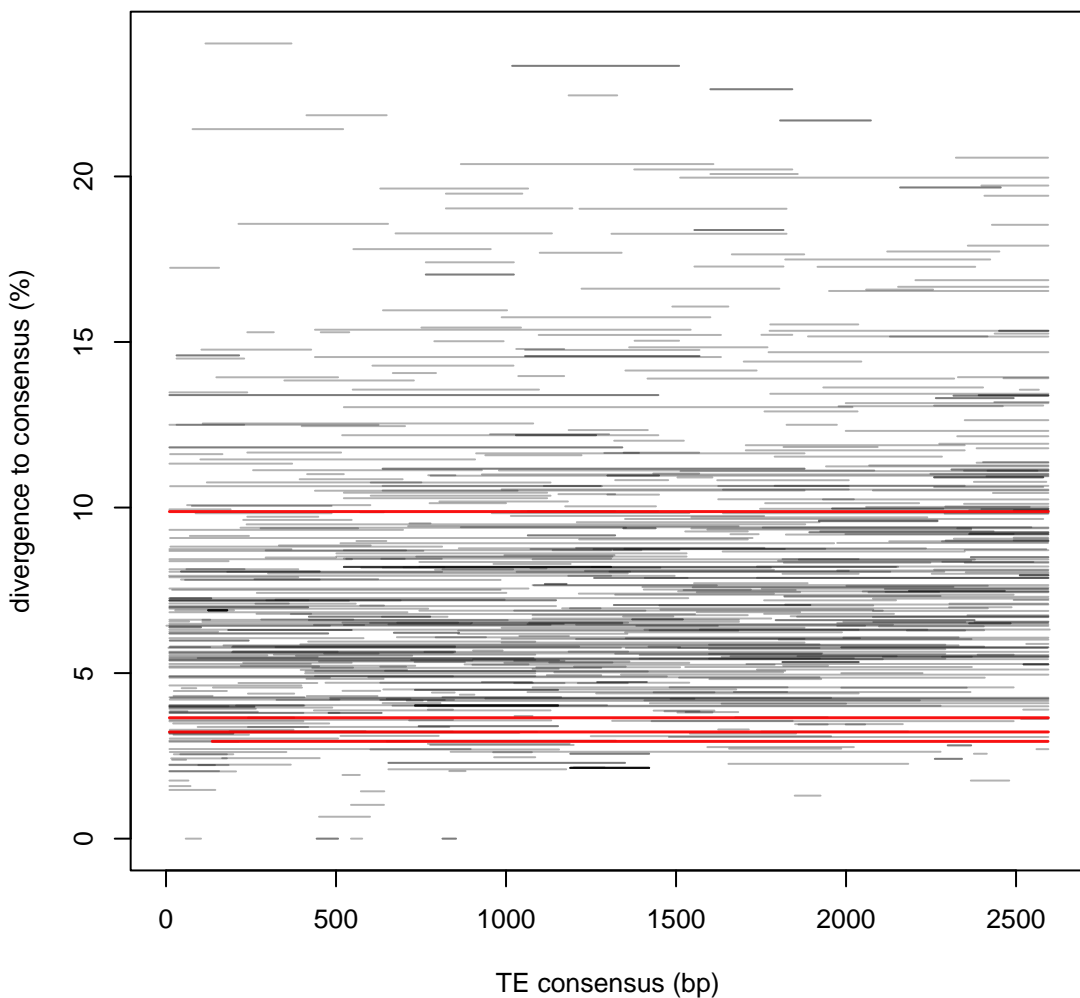
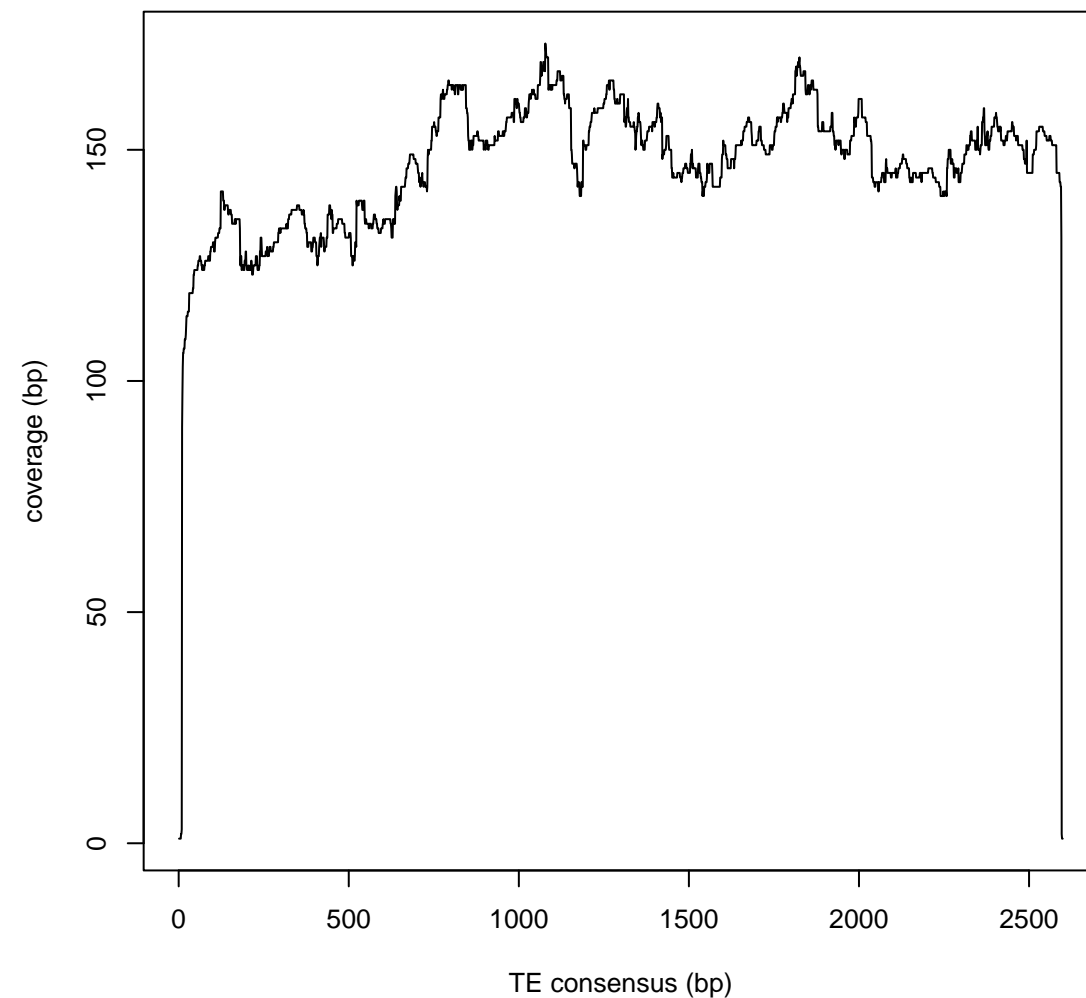


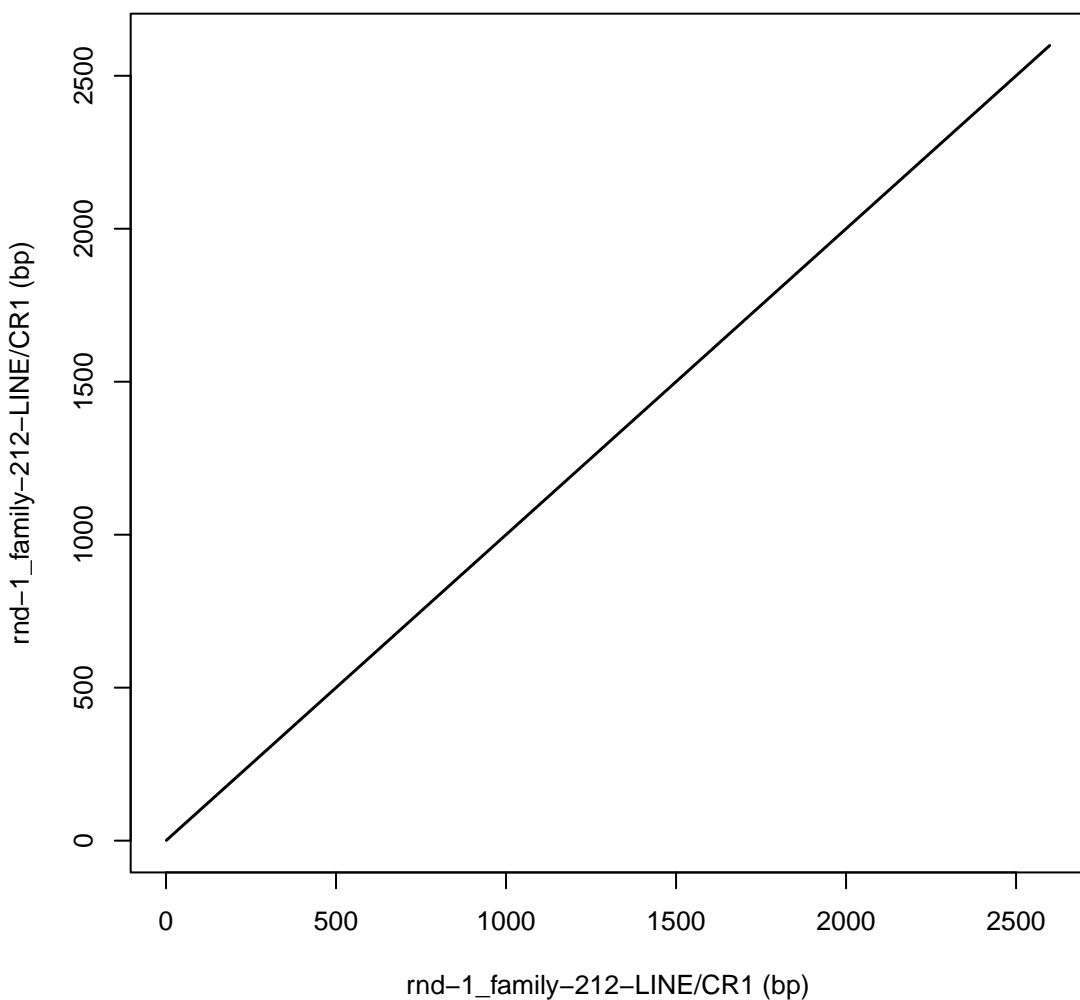
TE: rnd-1_family-212-LINE/CR1
consensus size: 2599bp; fragments: 739; full length: 4 (≥ 2339.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

