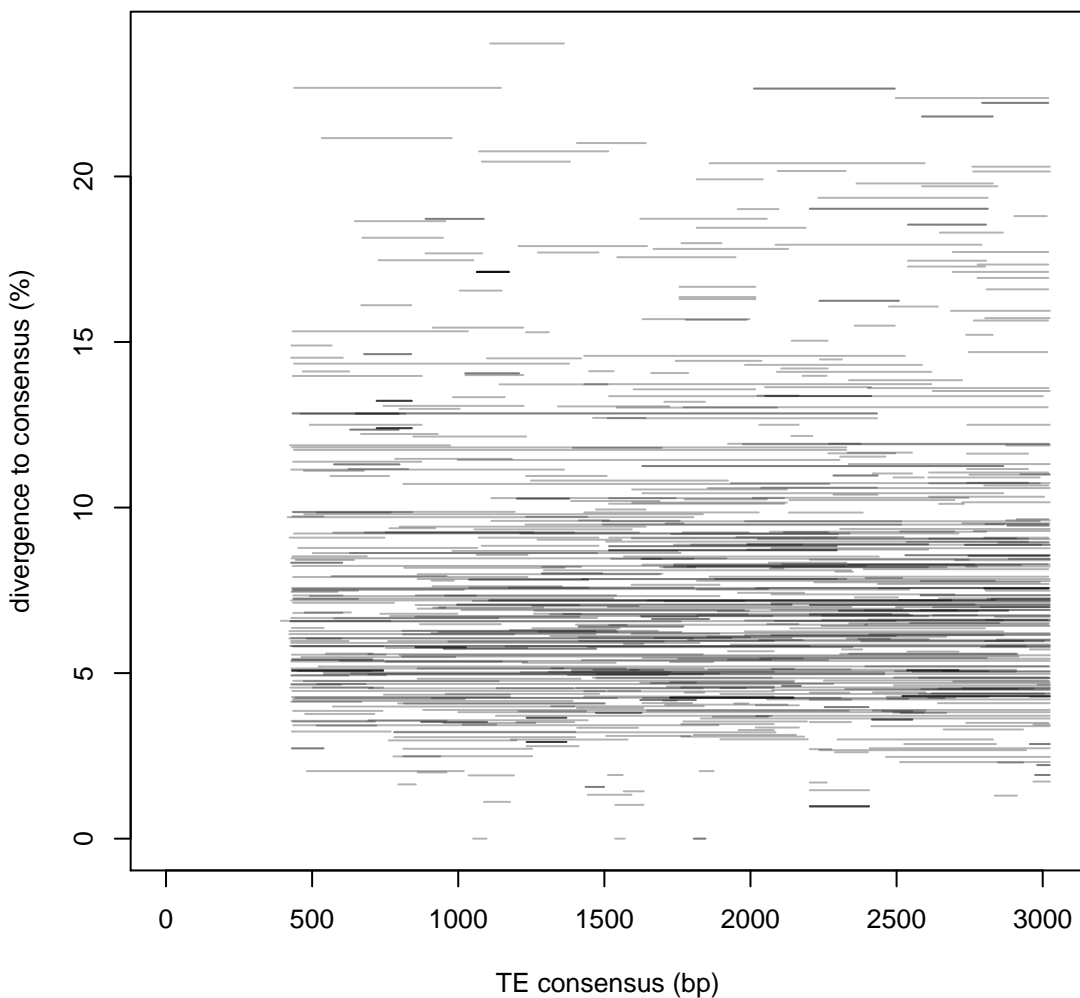
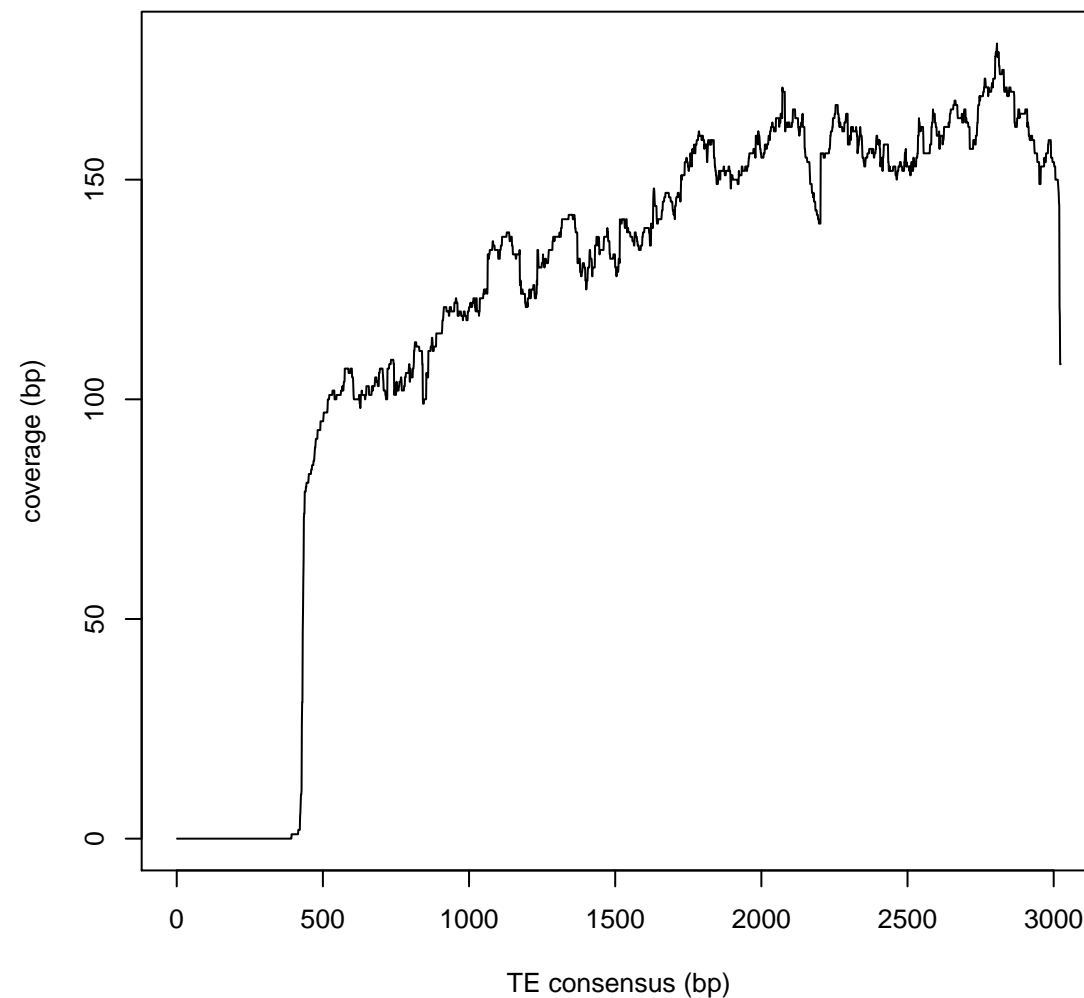


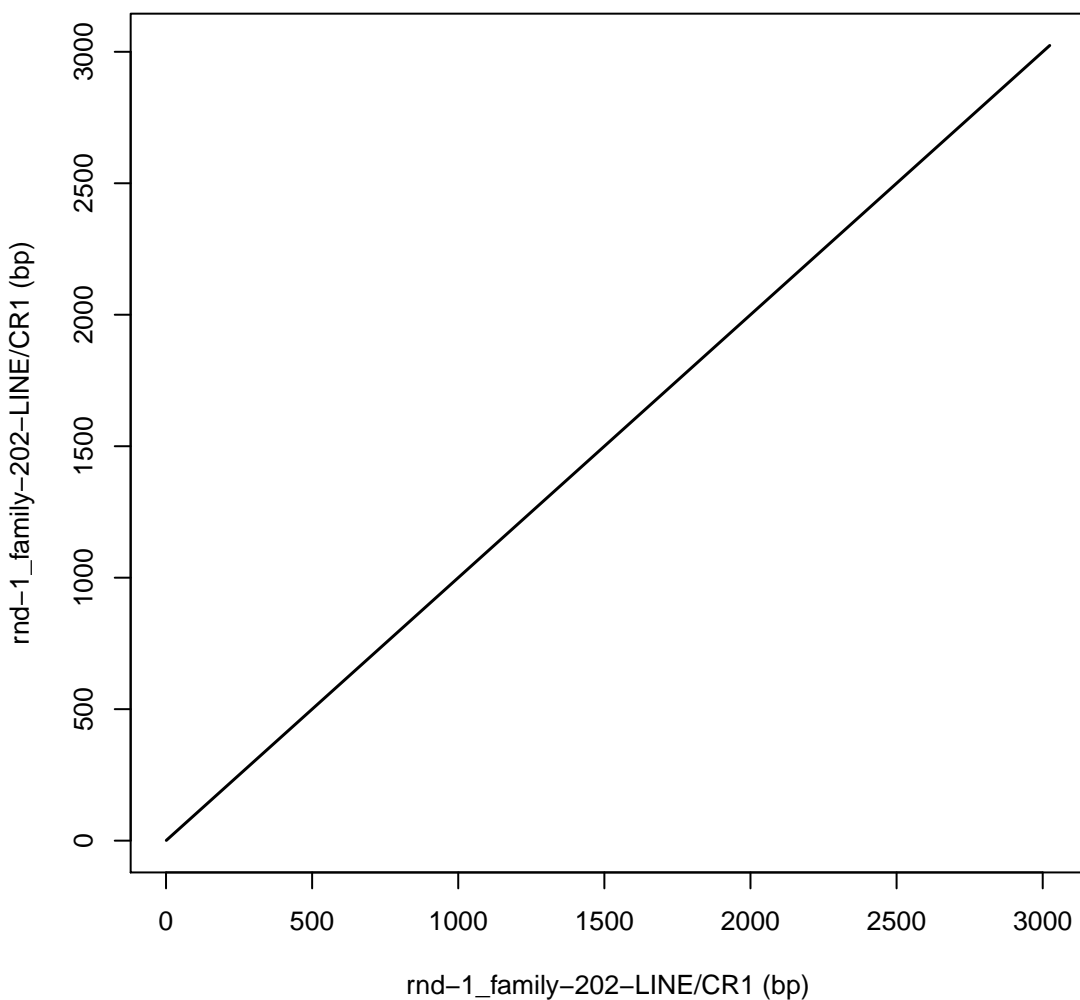
TE: rnd-1_family-202-LINE/CR1
consensus size: 3024bp; fragments: 704; full length: 0 (>=2721.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

