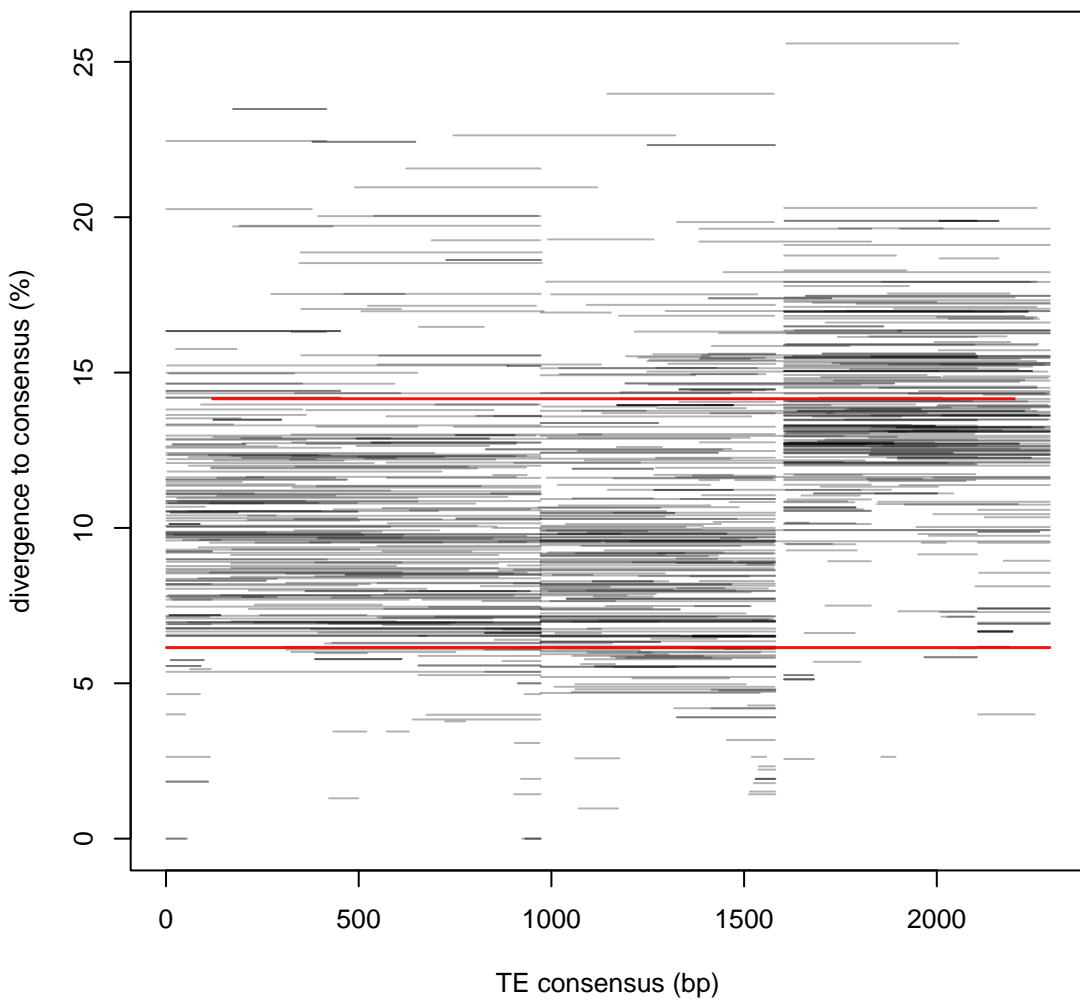
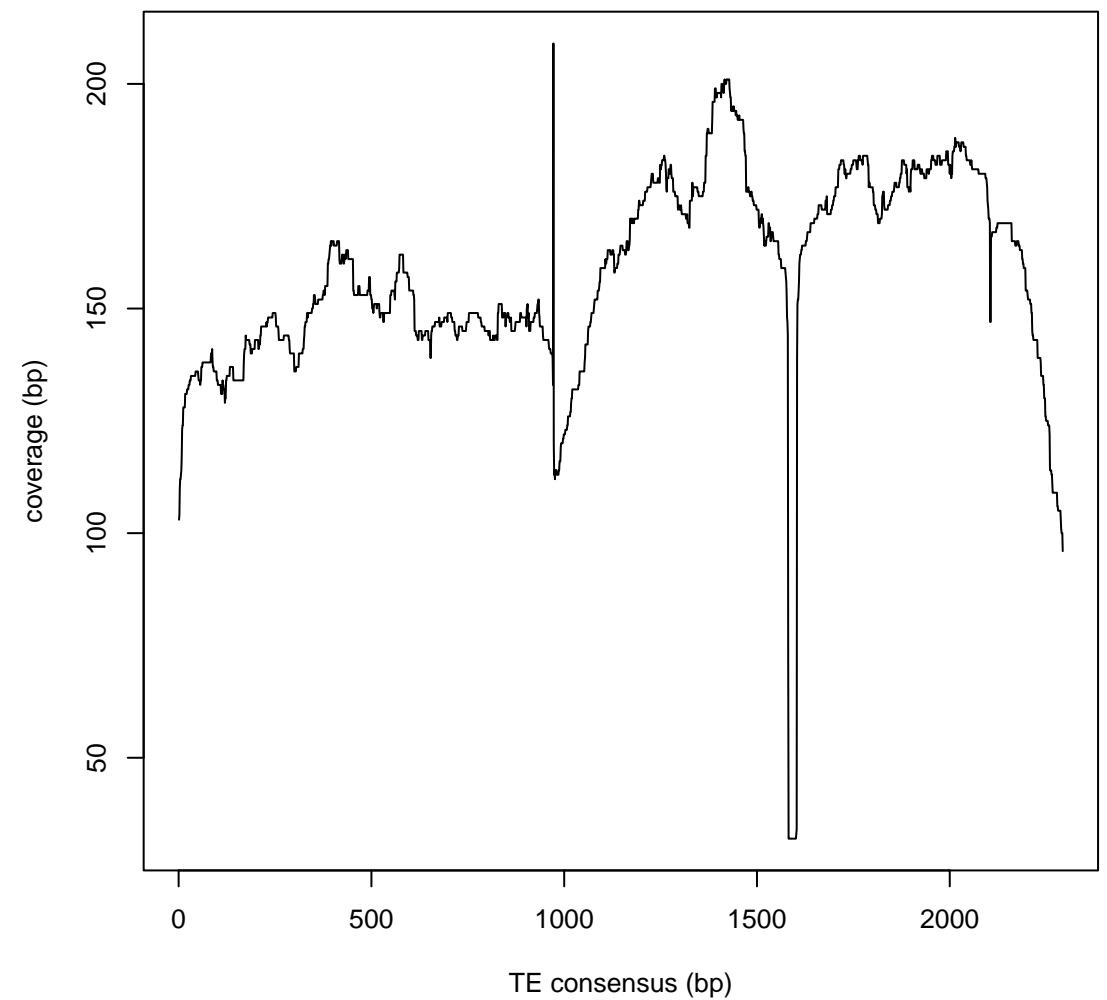


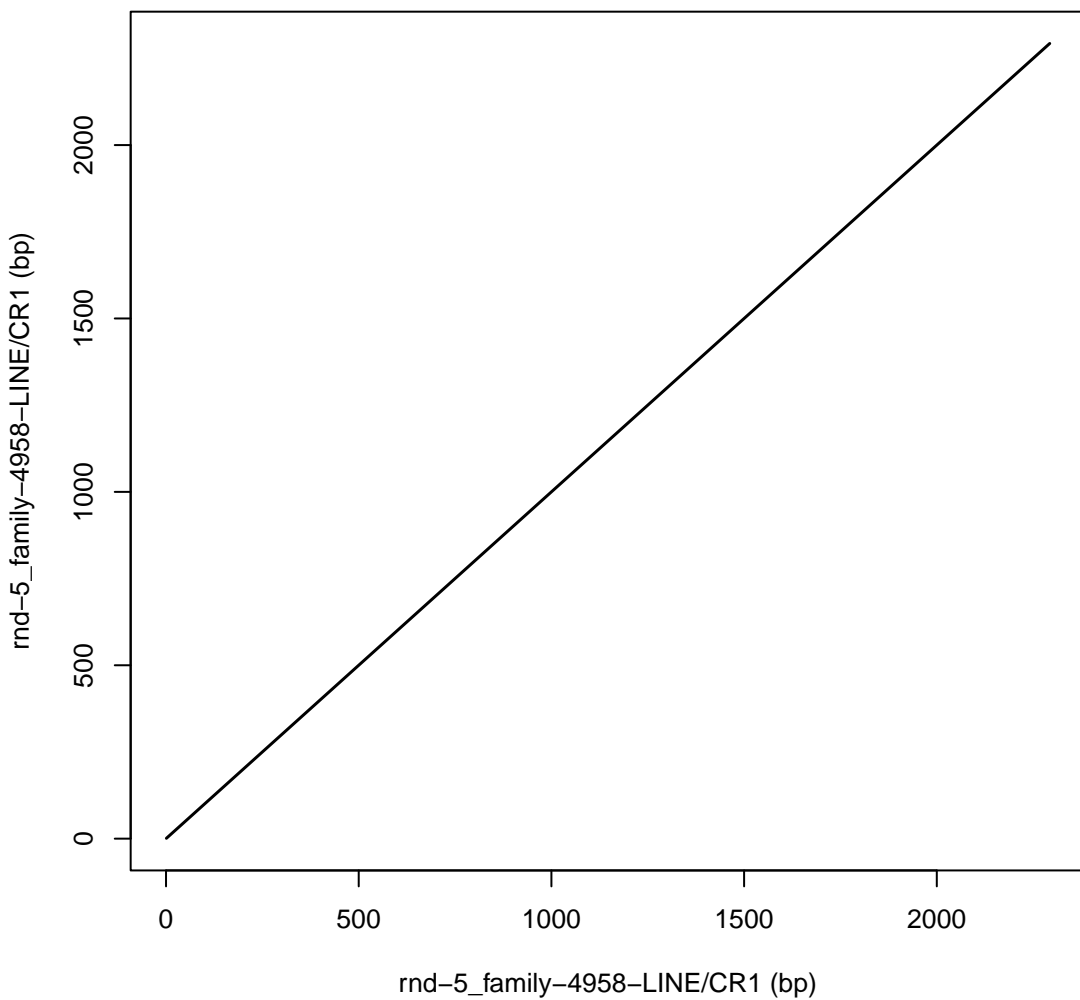
TE: rnd-5\_family-4958-LINE/CR1  
consensus size: 2293bp; fragments: 854; full length: 2 ( $\geq 2063.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

