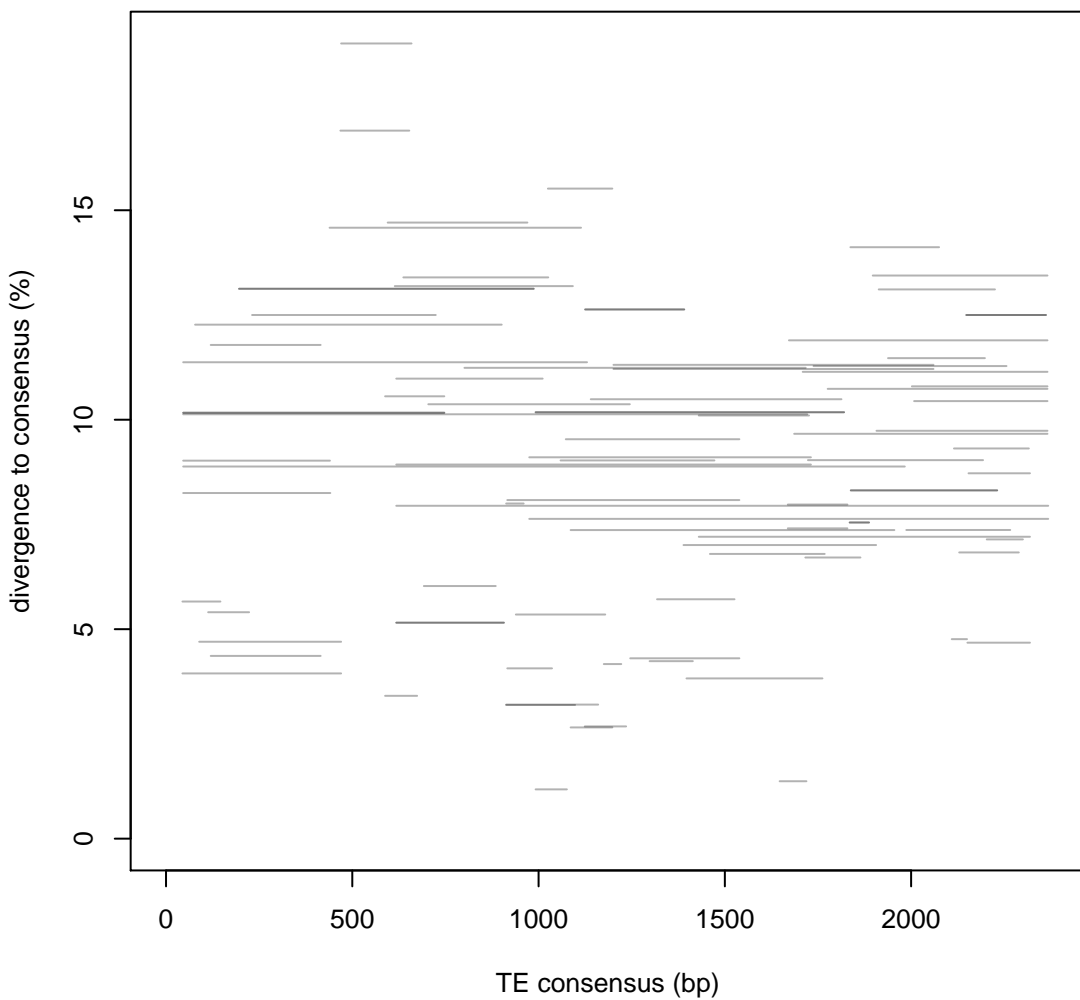
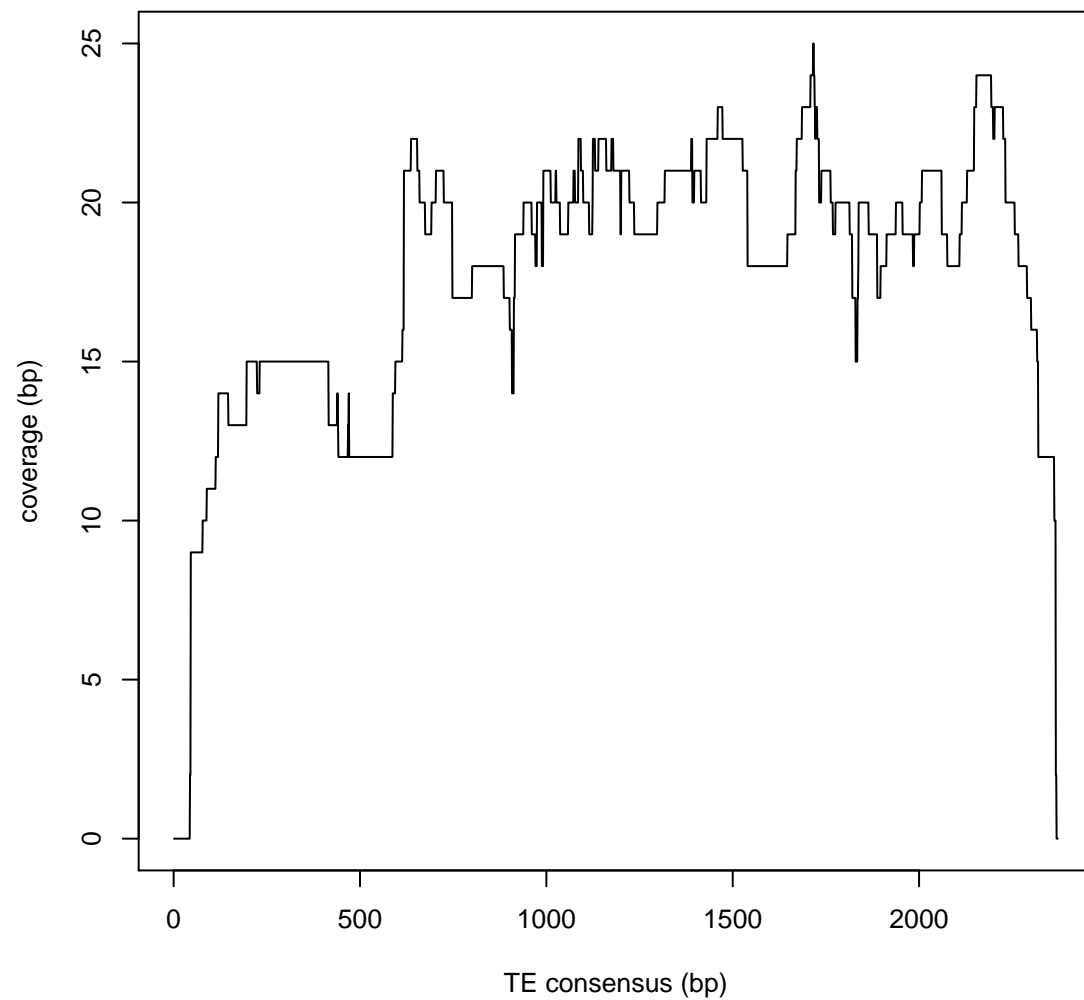


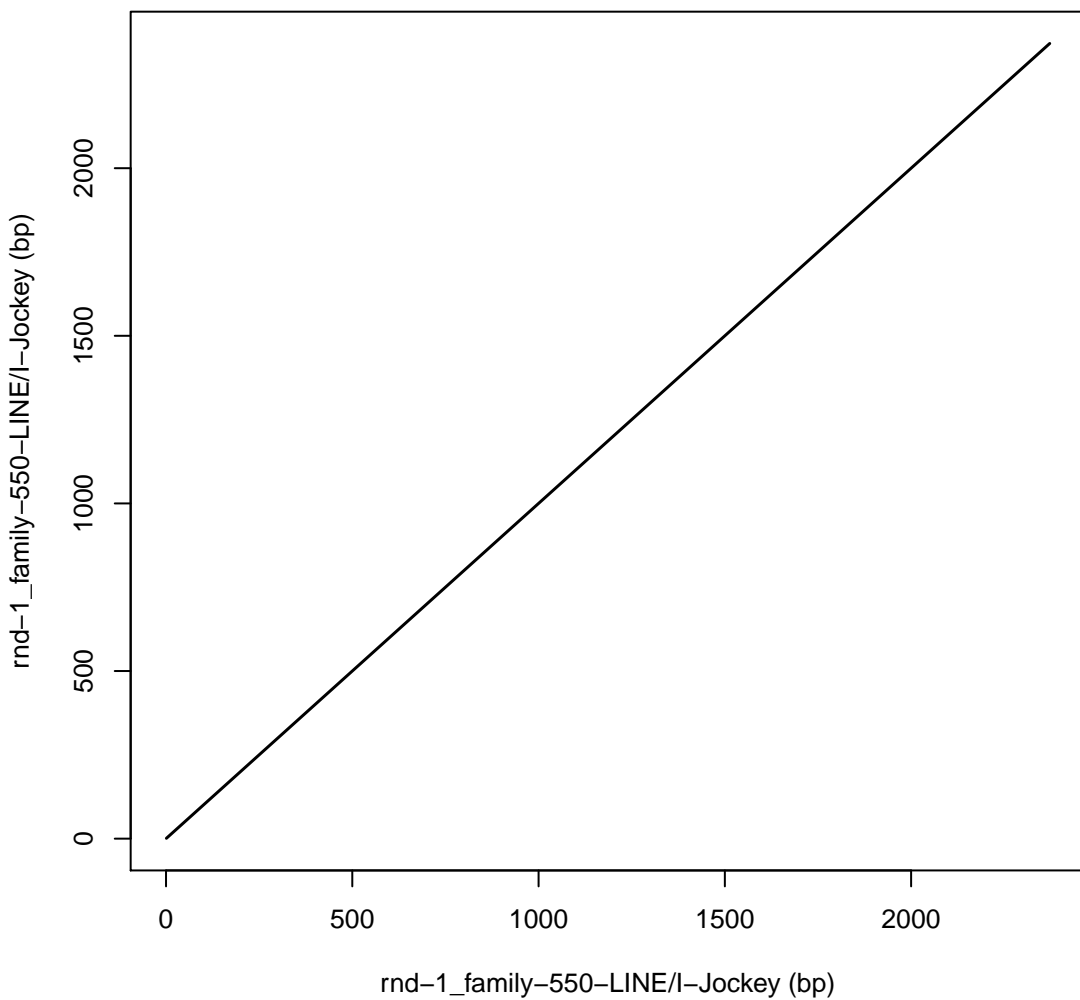
TE: rnd-1\_family-550-LINE/I-Jockey  
consensus size: 2372bp; fragments: 94; full length: 0 ( $\geq 2134.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

