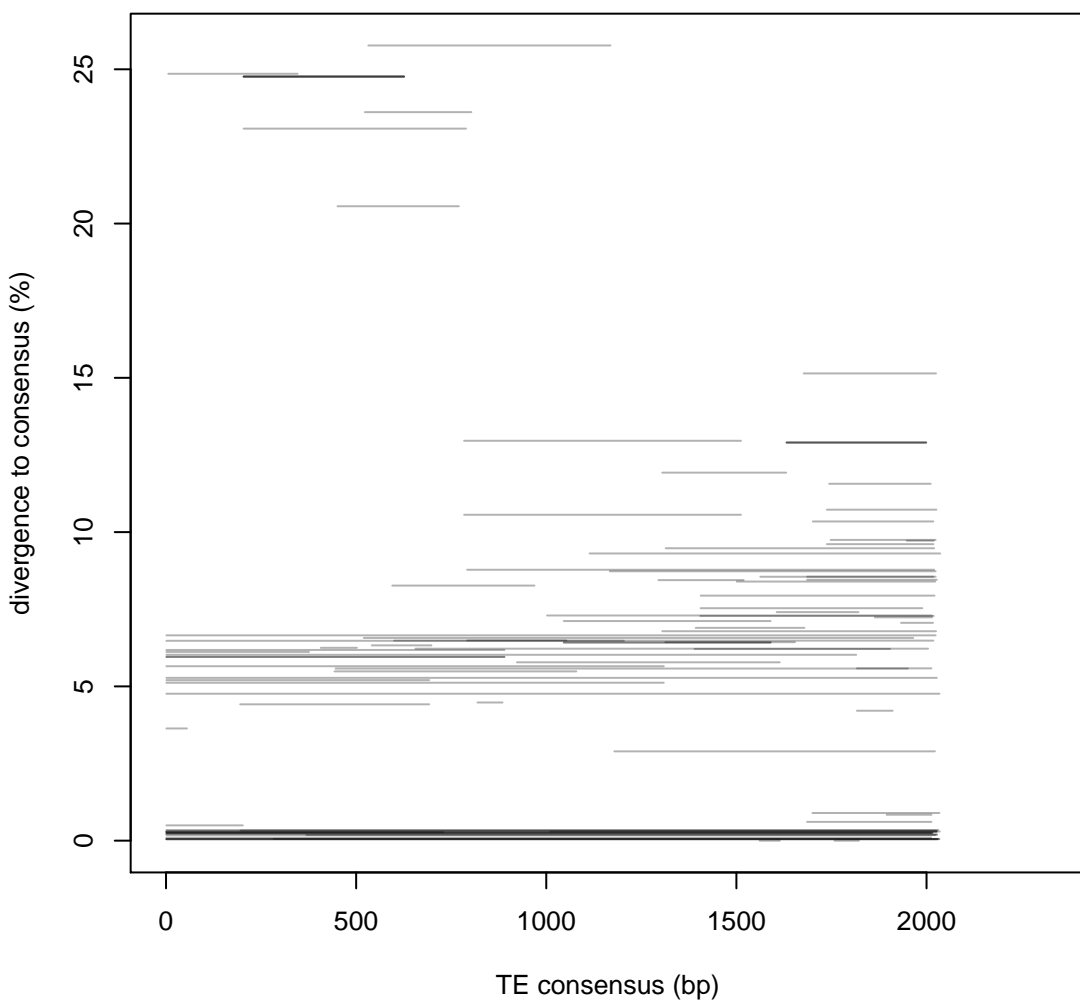
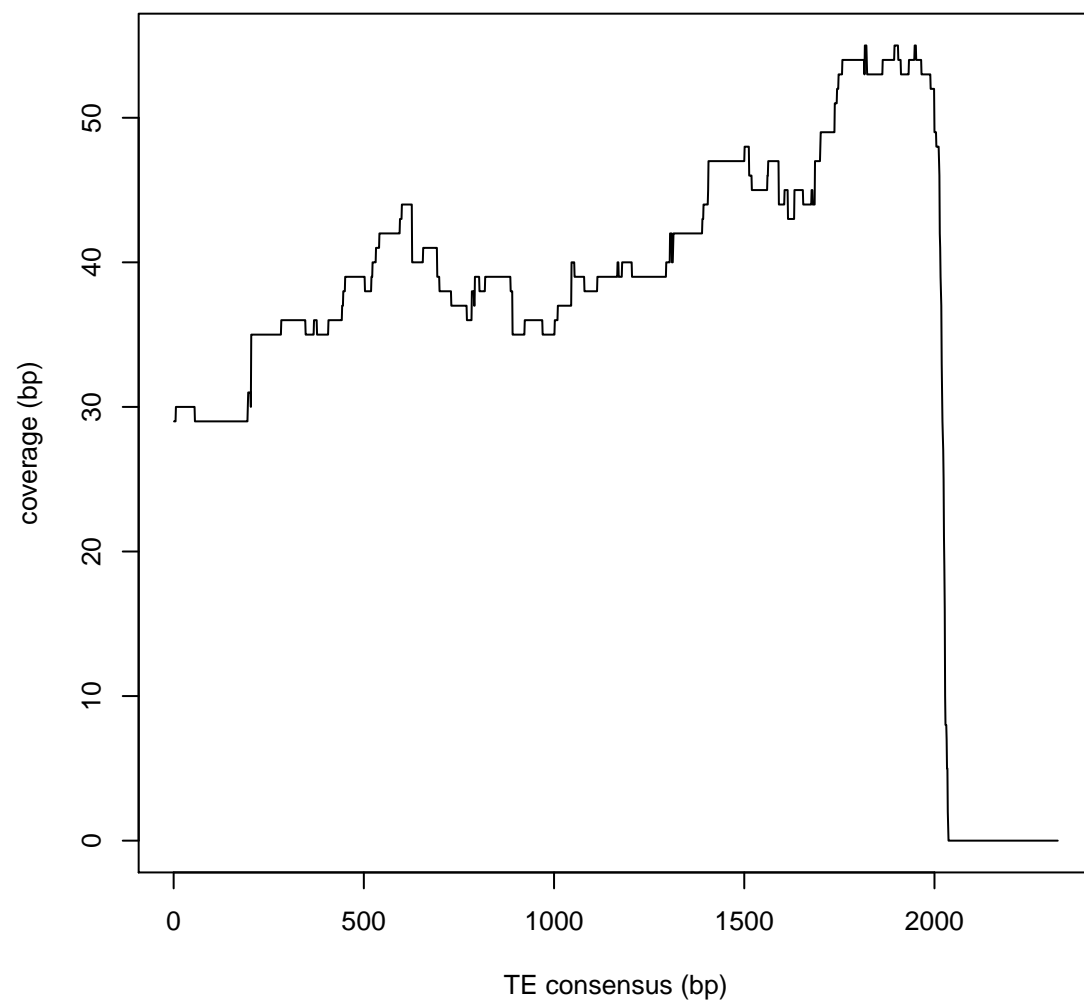


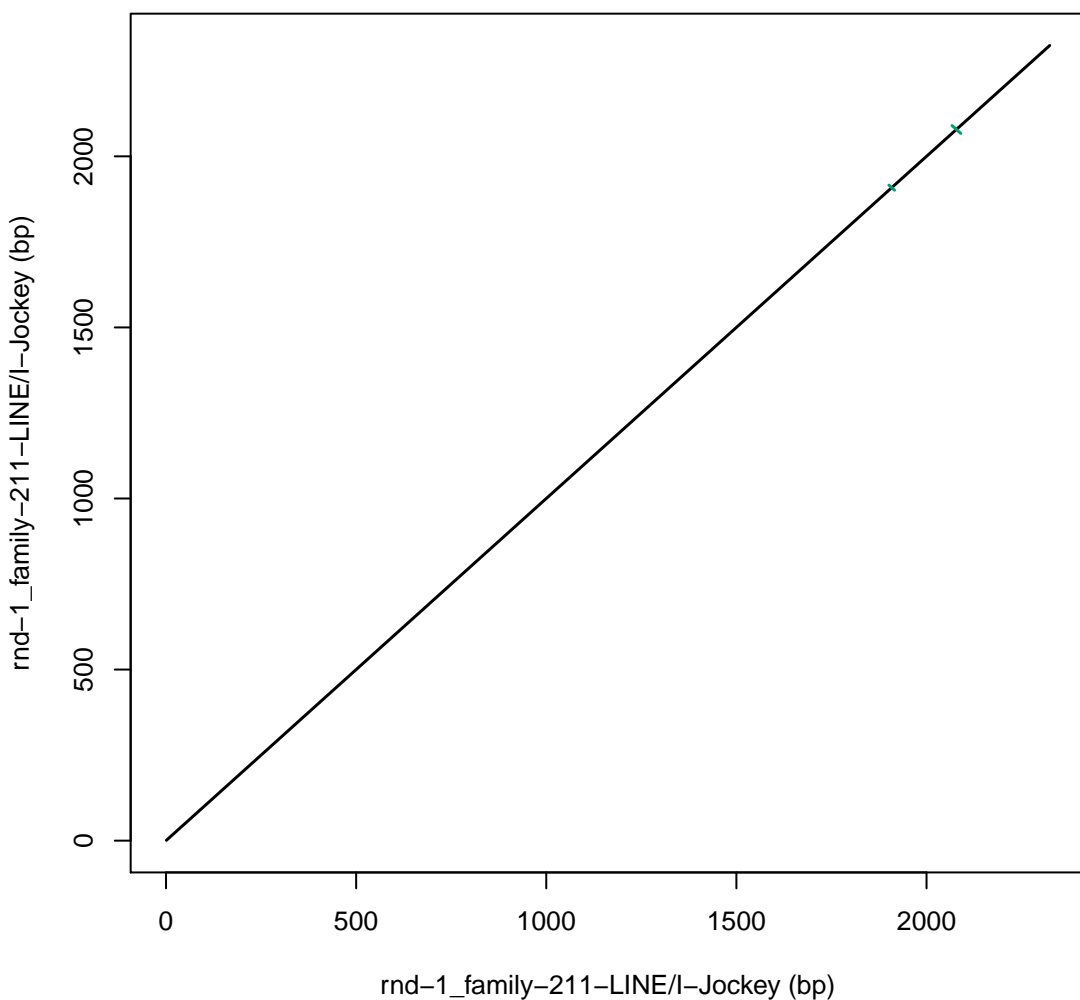
TE: rnd-1\_family-211-LINE/I-Jockey  
consensus size: 2324bp; fragments: 98; full length: 0 ( $\geq 2091.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

