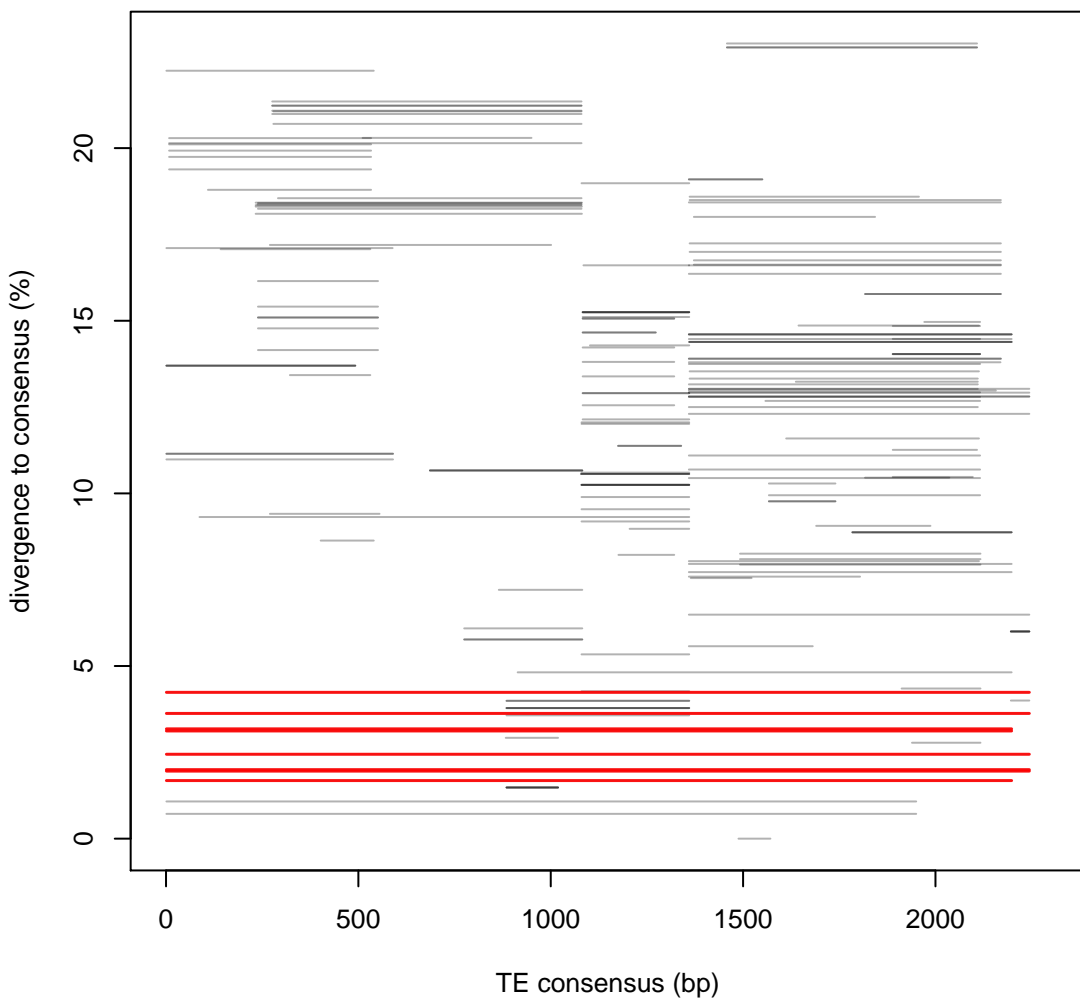
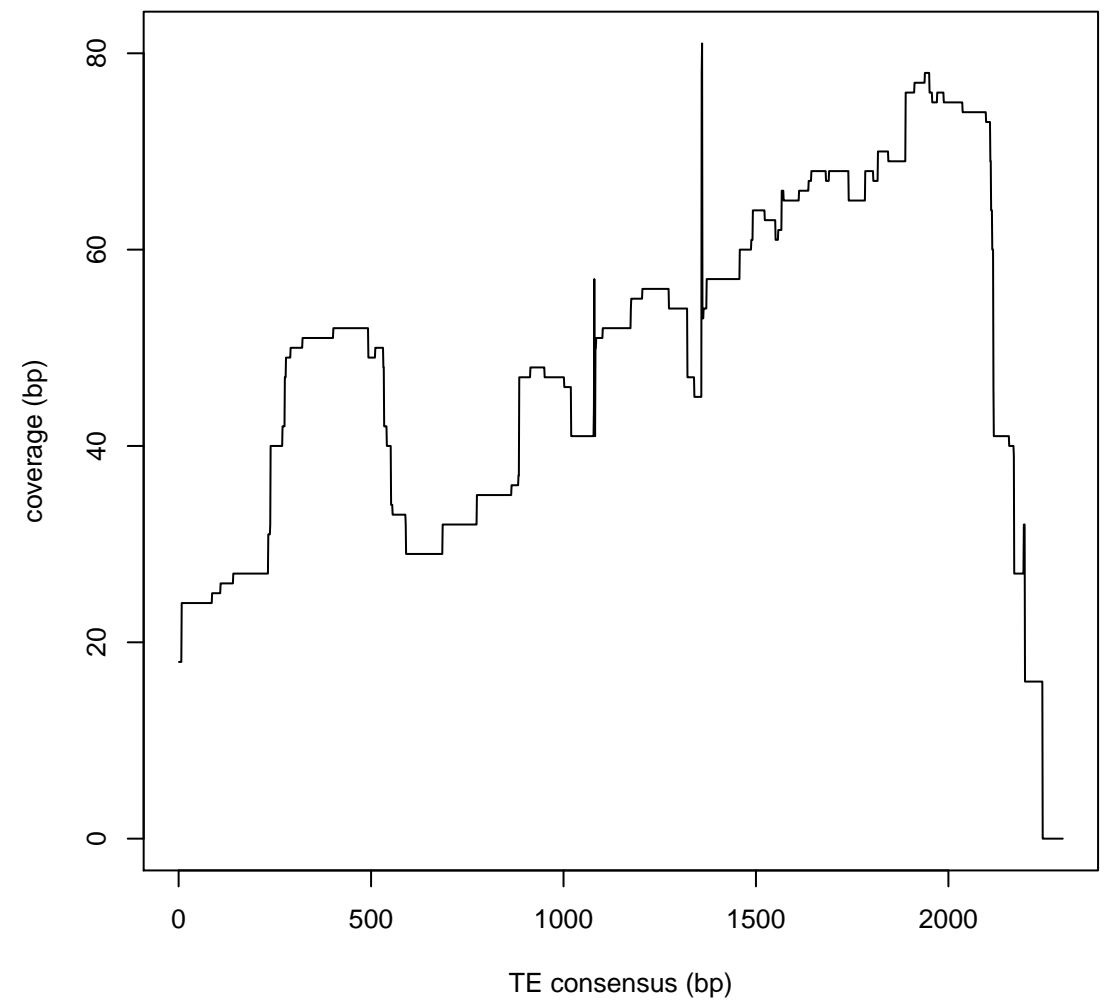


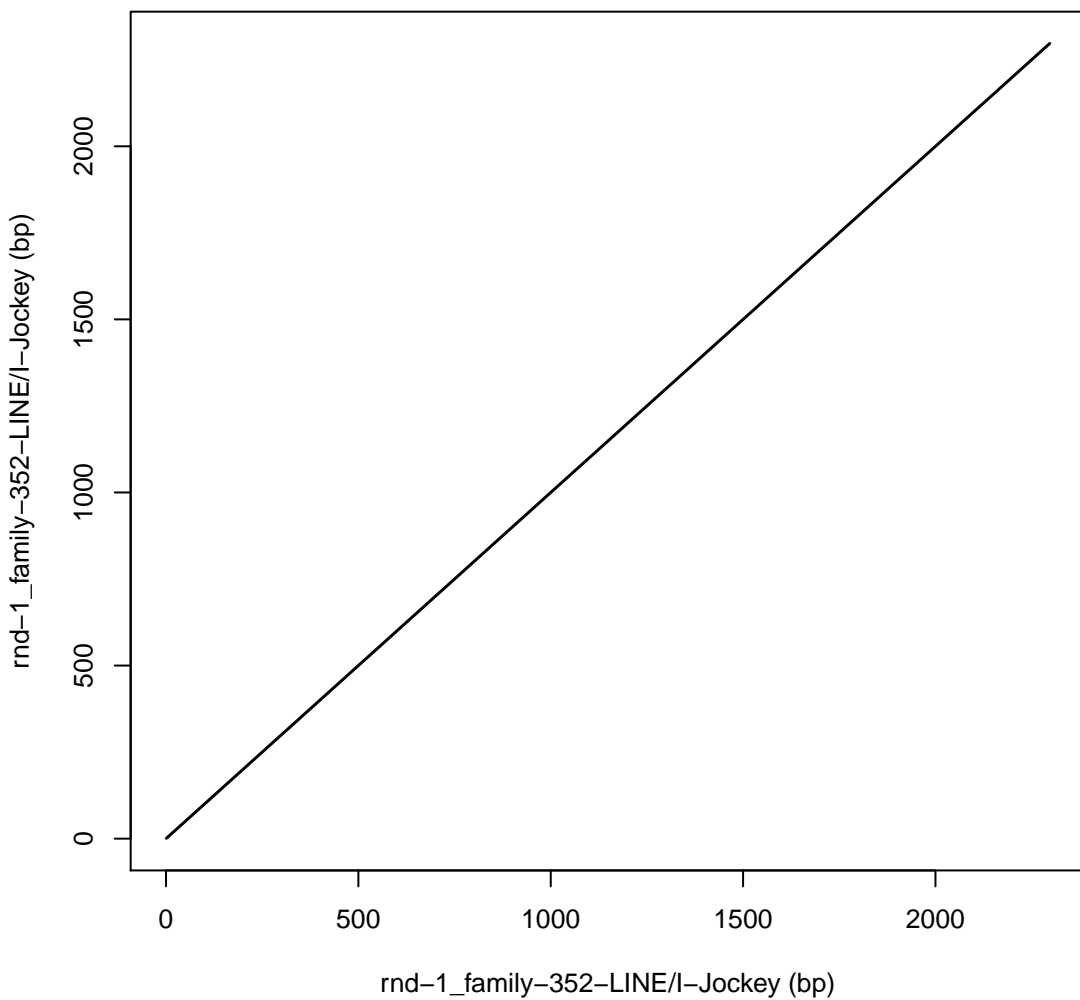
TE: rnd-1\_family-352-LINE/I-Jockey  
consensus size: 2297bp; fragments: 193; full length: 8 ( $\geq 2067.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

