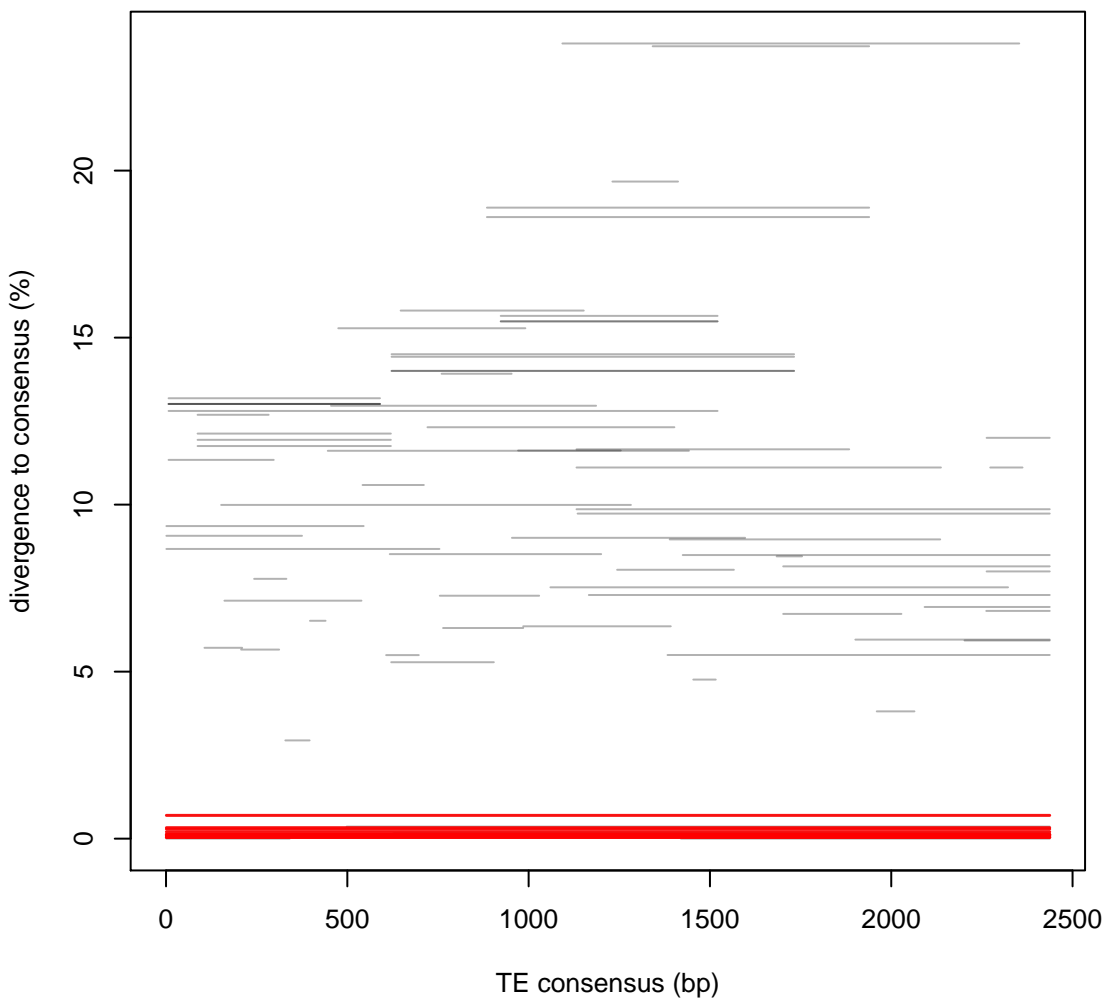
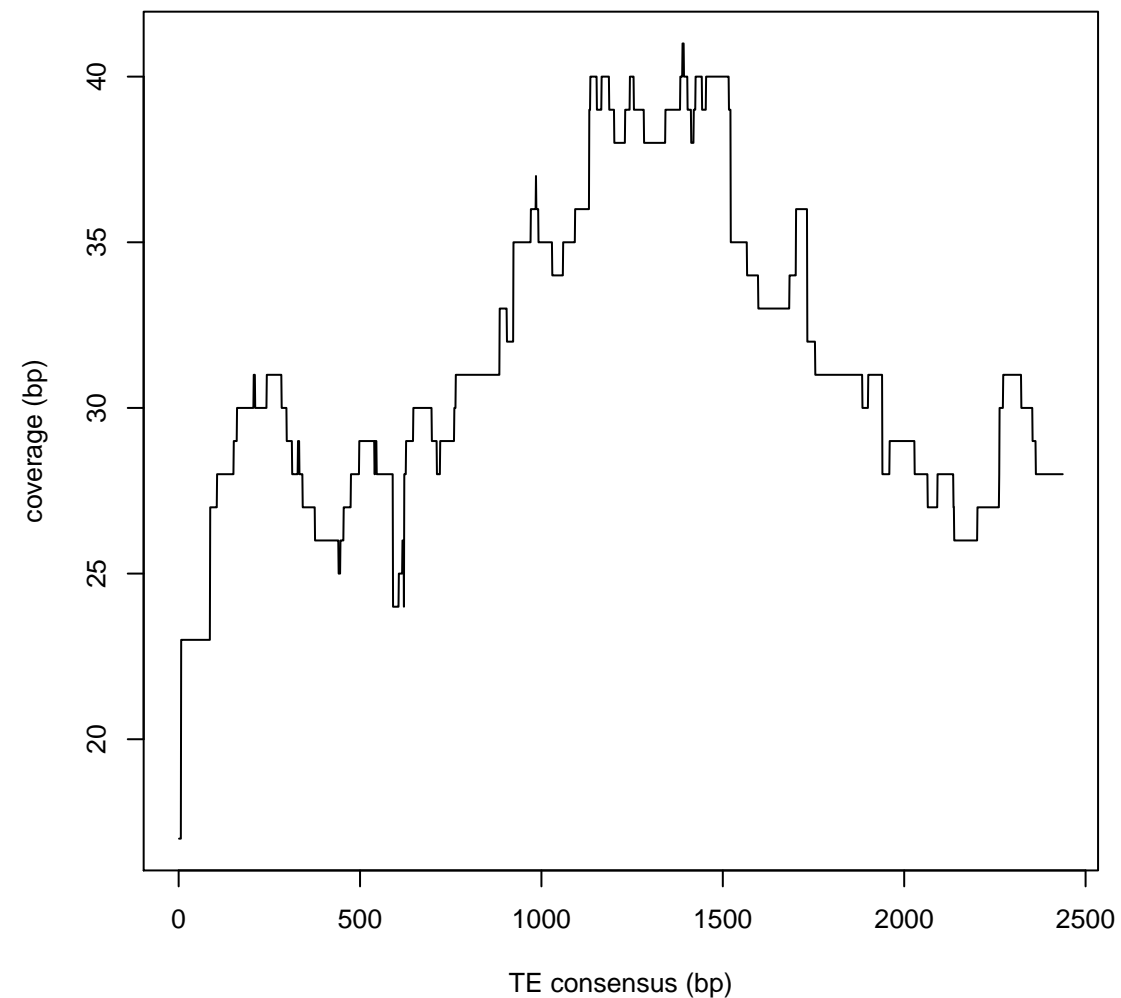


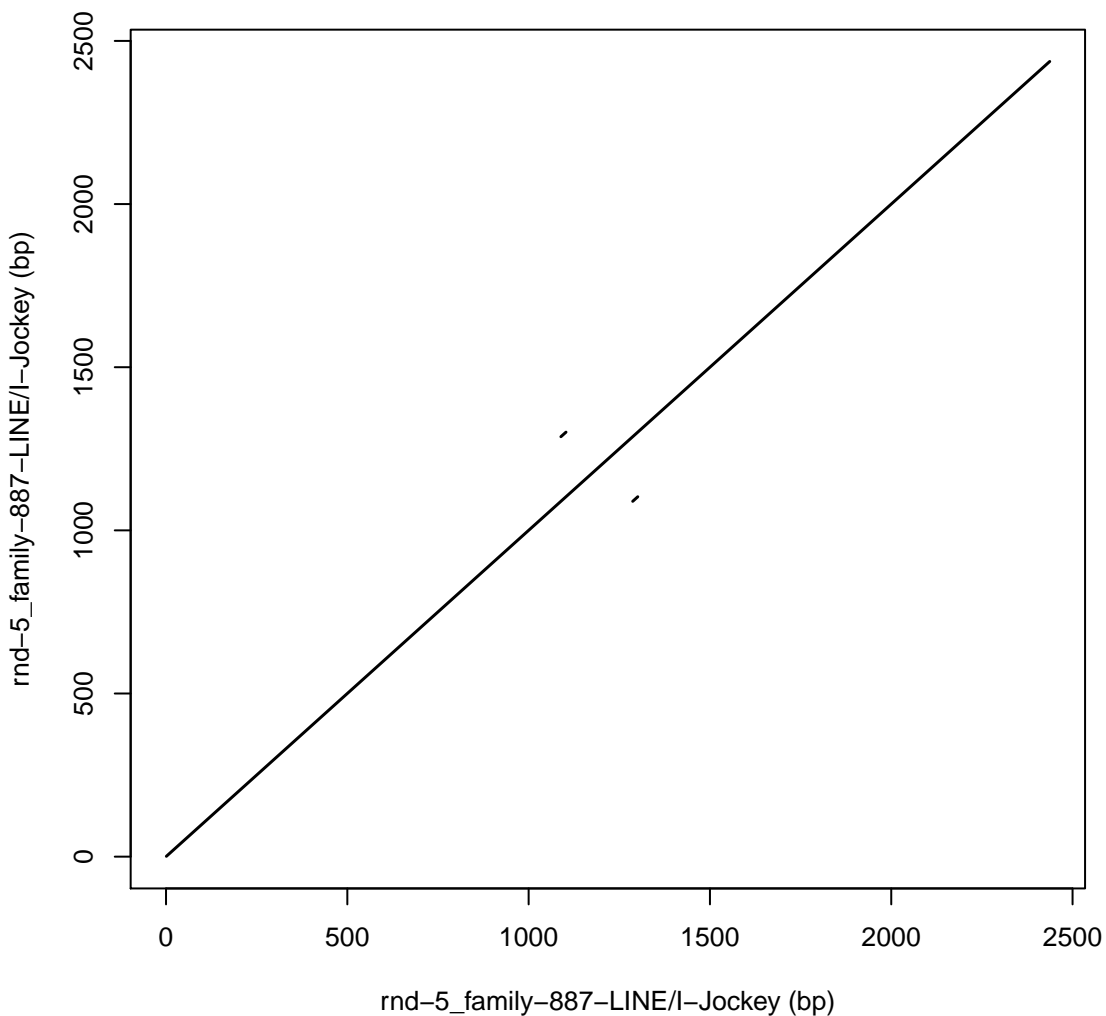
TE: rnd-5\_family-887-LINE/I-Jockey  
consensus size: 2437bp; fragments: 86; full length: 13 ( $\geq 2193.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

