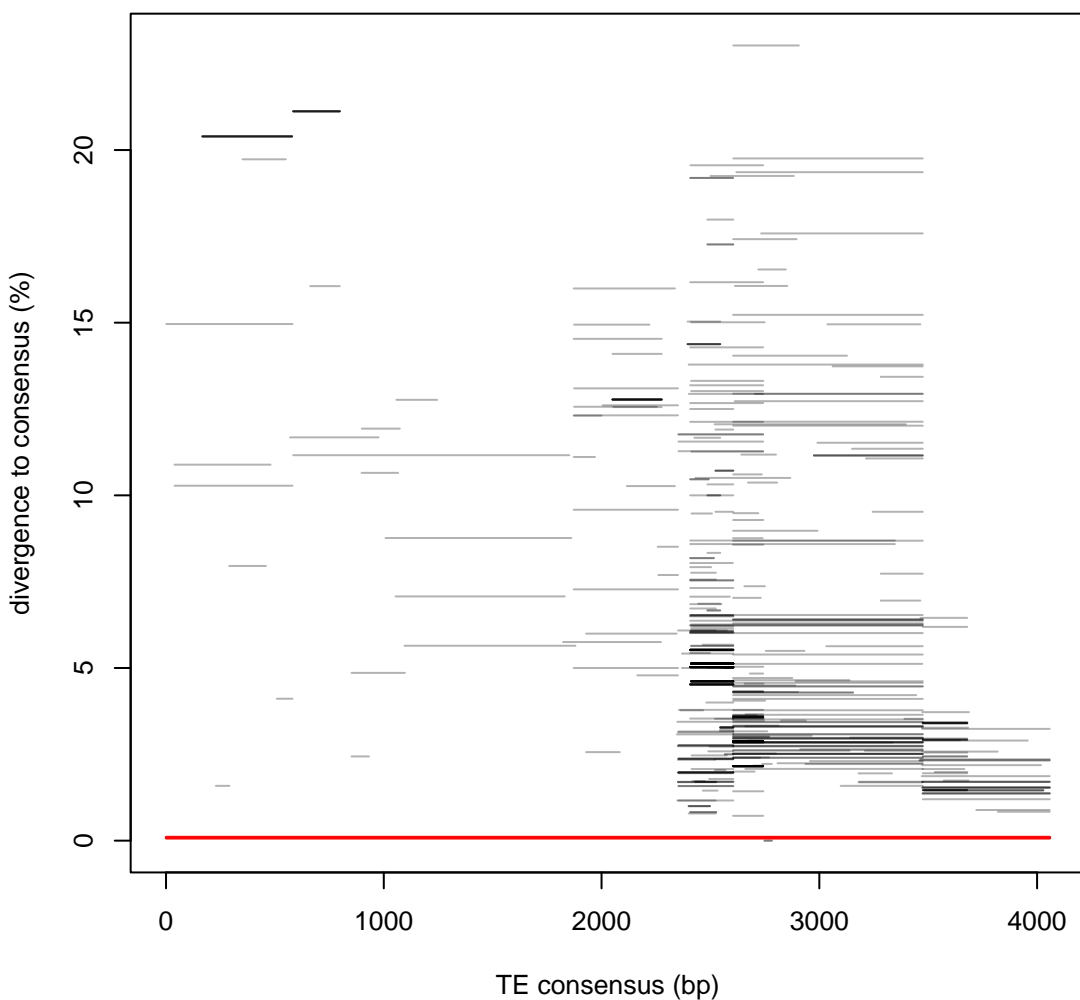
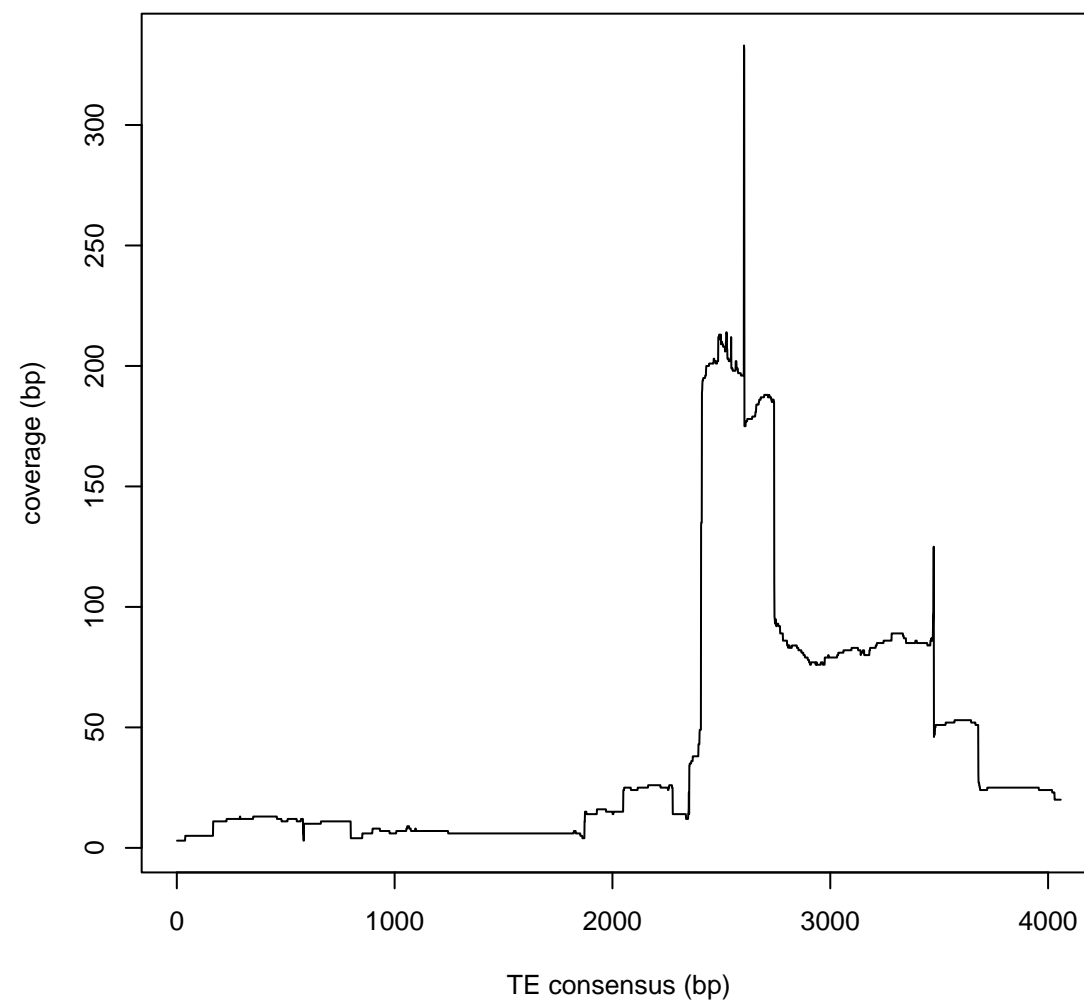


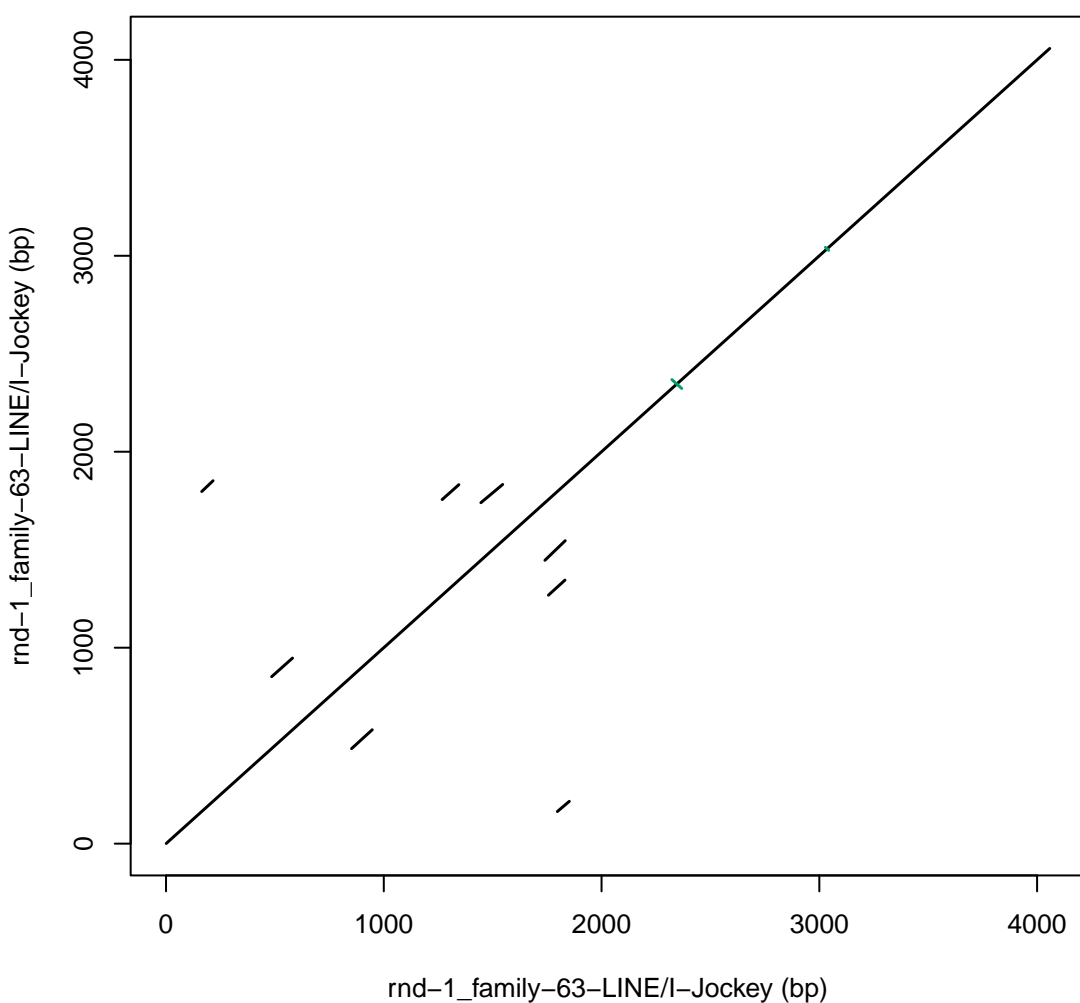
TE: rnd-1\_family-63-LINE/I-Jockey  
consensus size: 4058bp; fragments: 543; full length: 2 ( $\geq 3652.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

