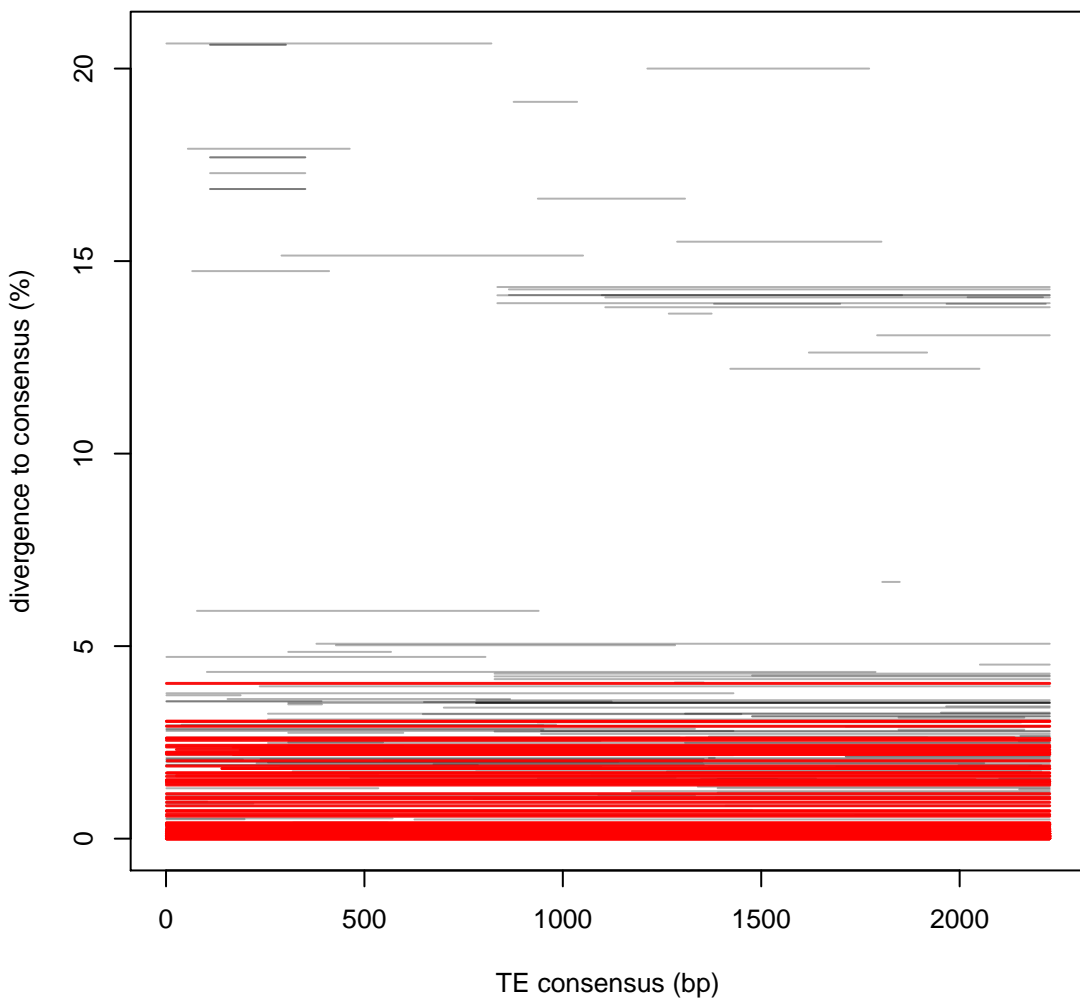
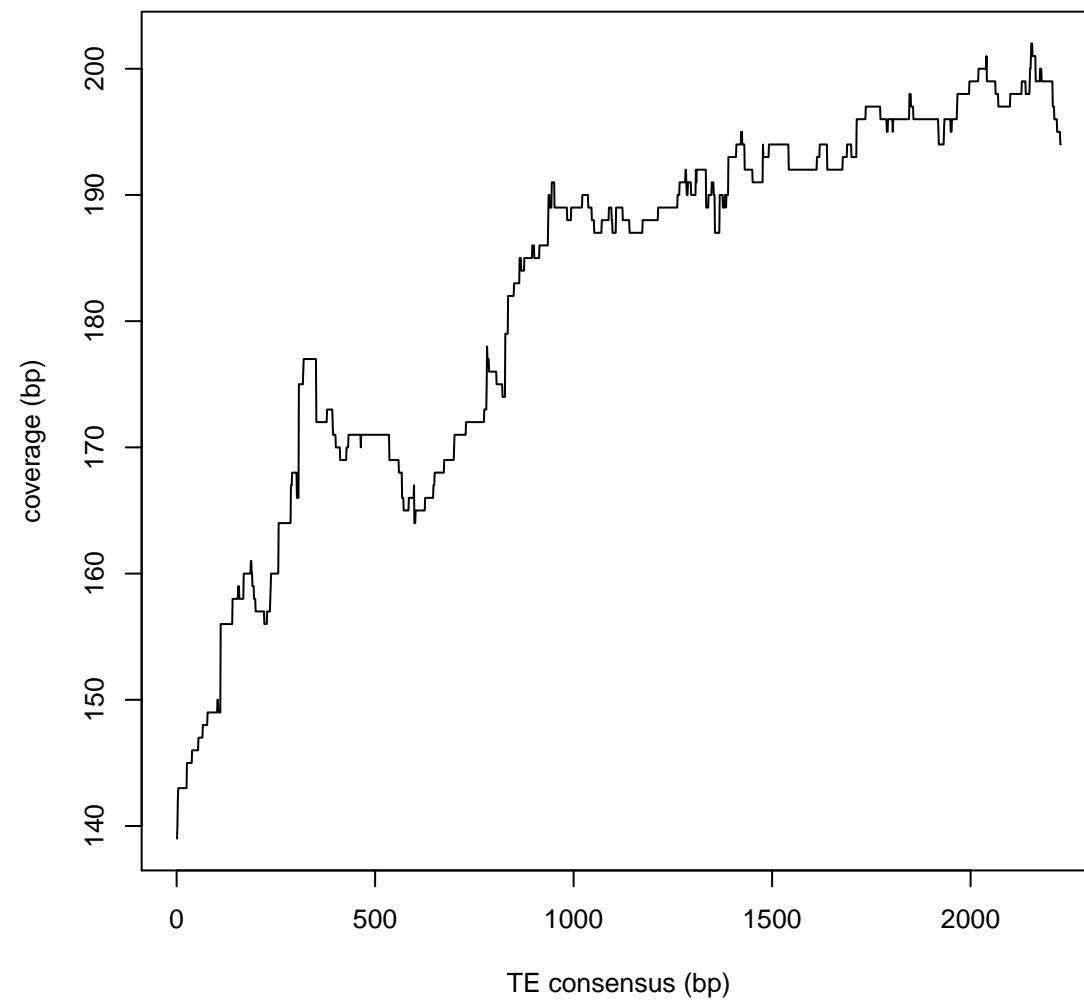


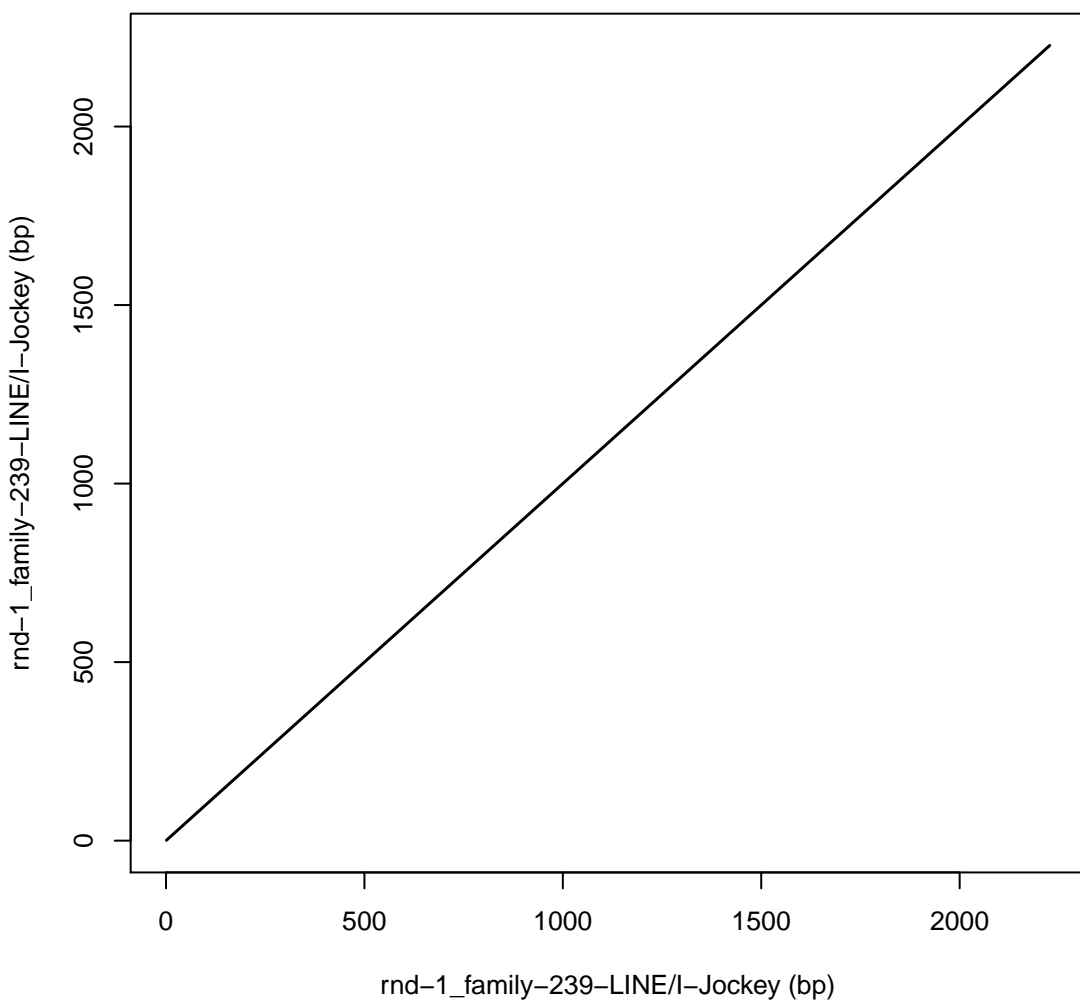
TE: rnd-1\_family-239-LINE/I-Jockey  
consensus size: 2227bp; fragments: 289; full length: 115 (>=2004.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

