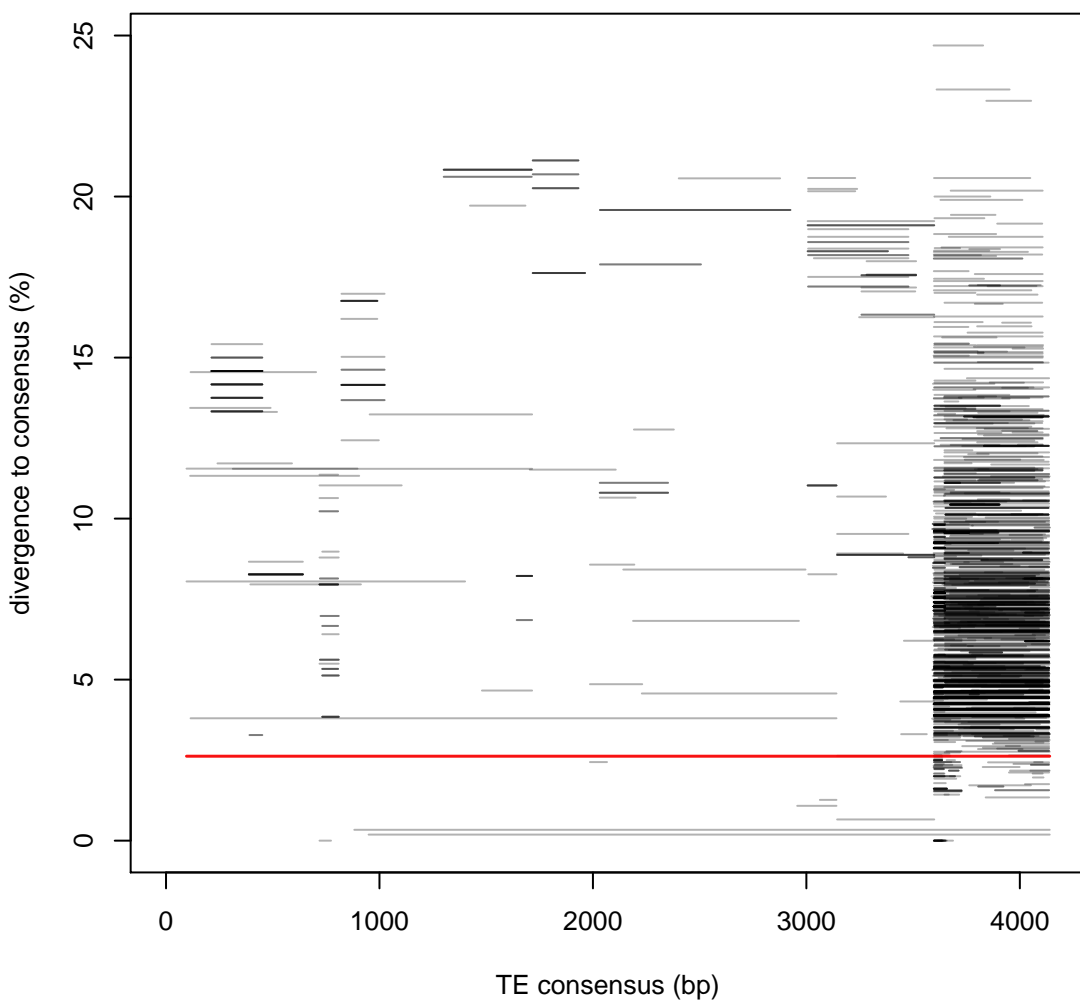
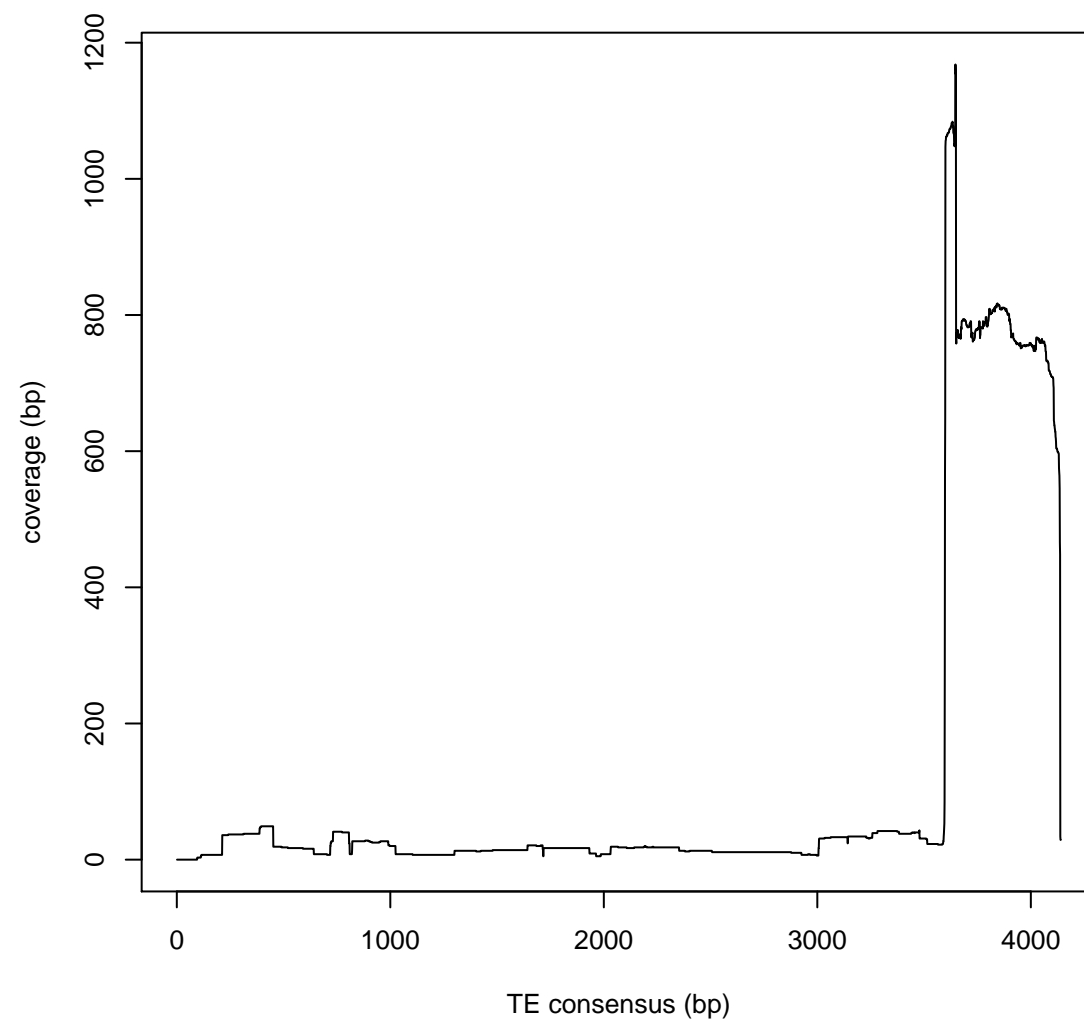


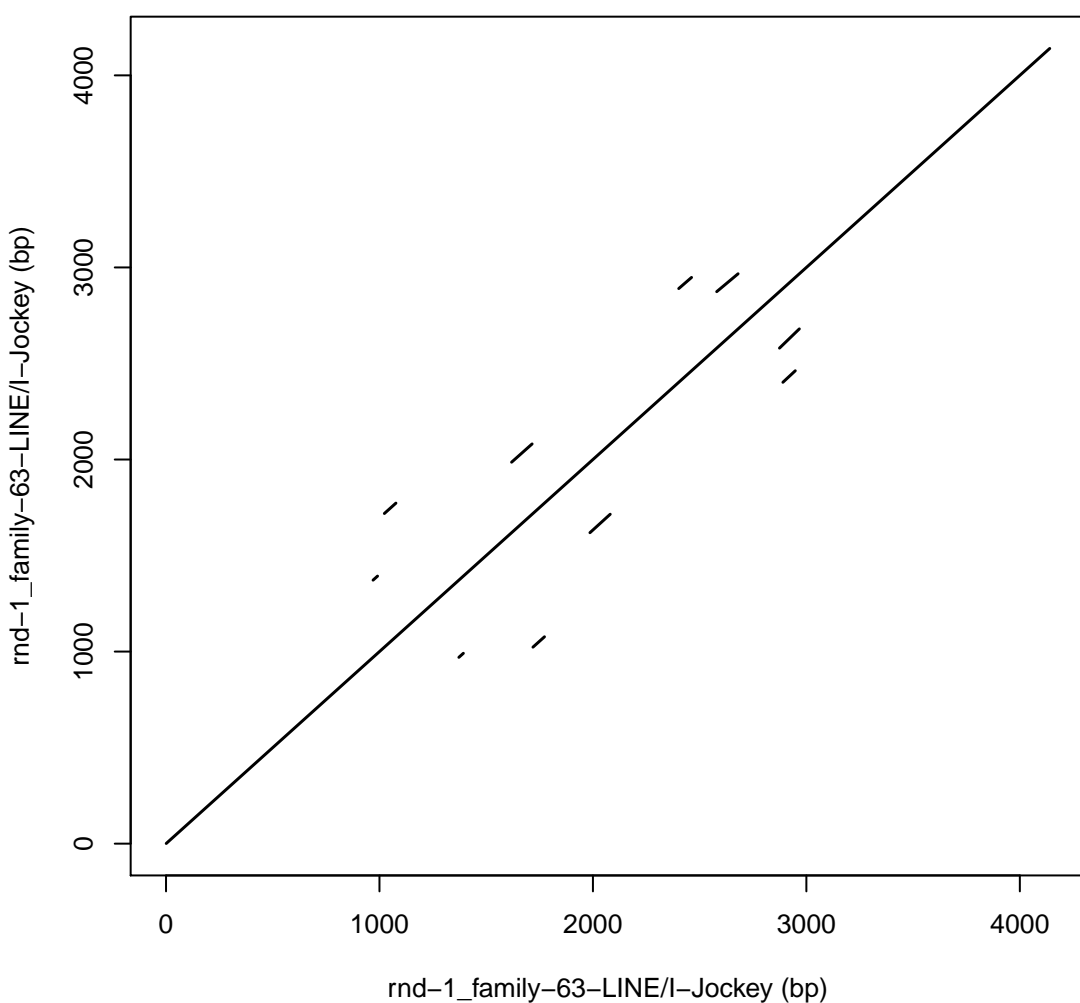
TE: rnd-1_family-63-LINE/I-Jockey
consensus size: 4140bp; fragments: 1825; full length: 1 (>=3726bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

