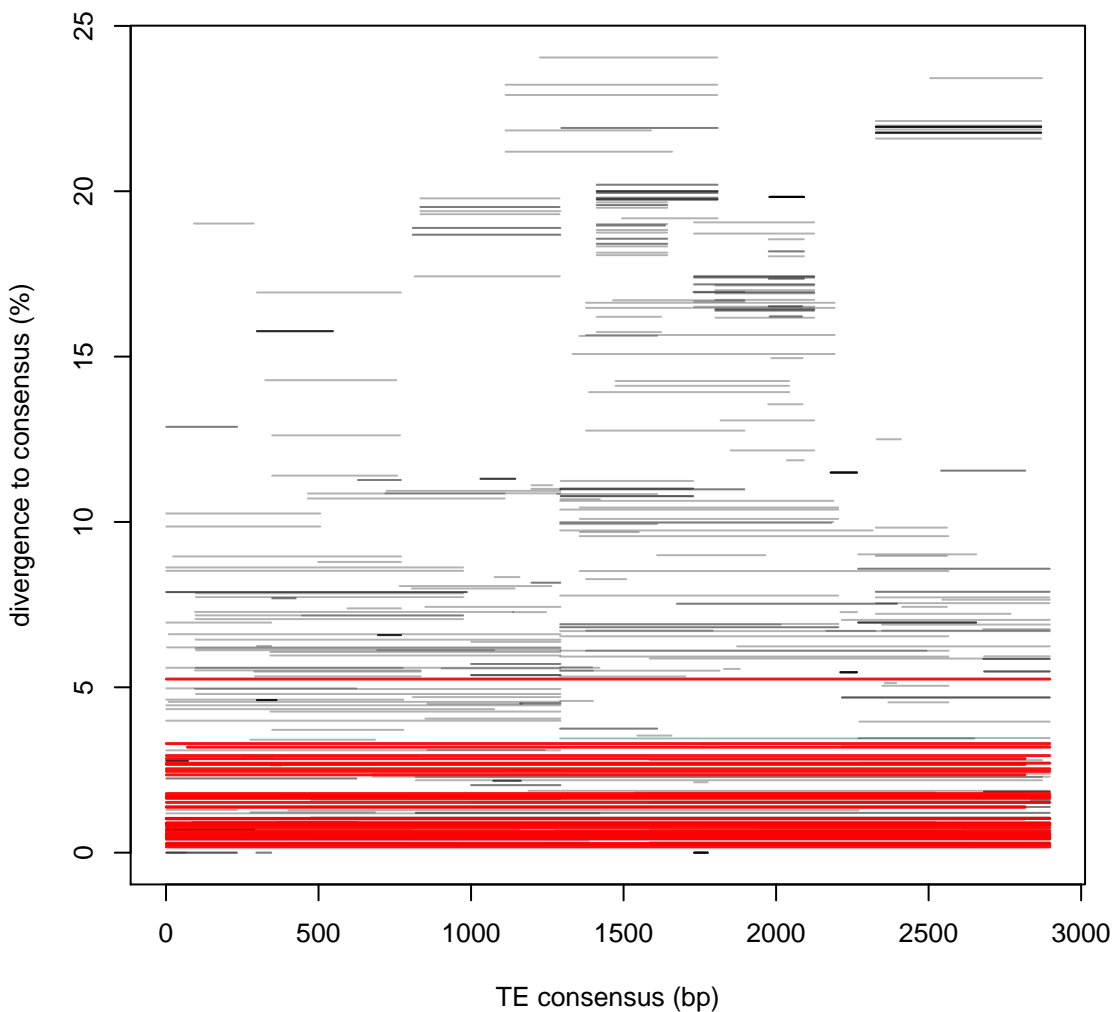
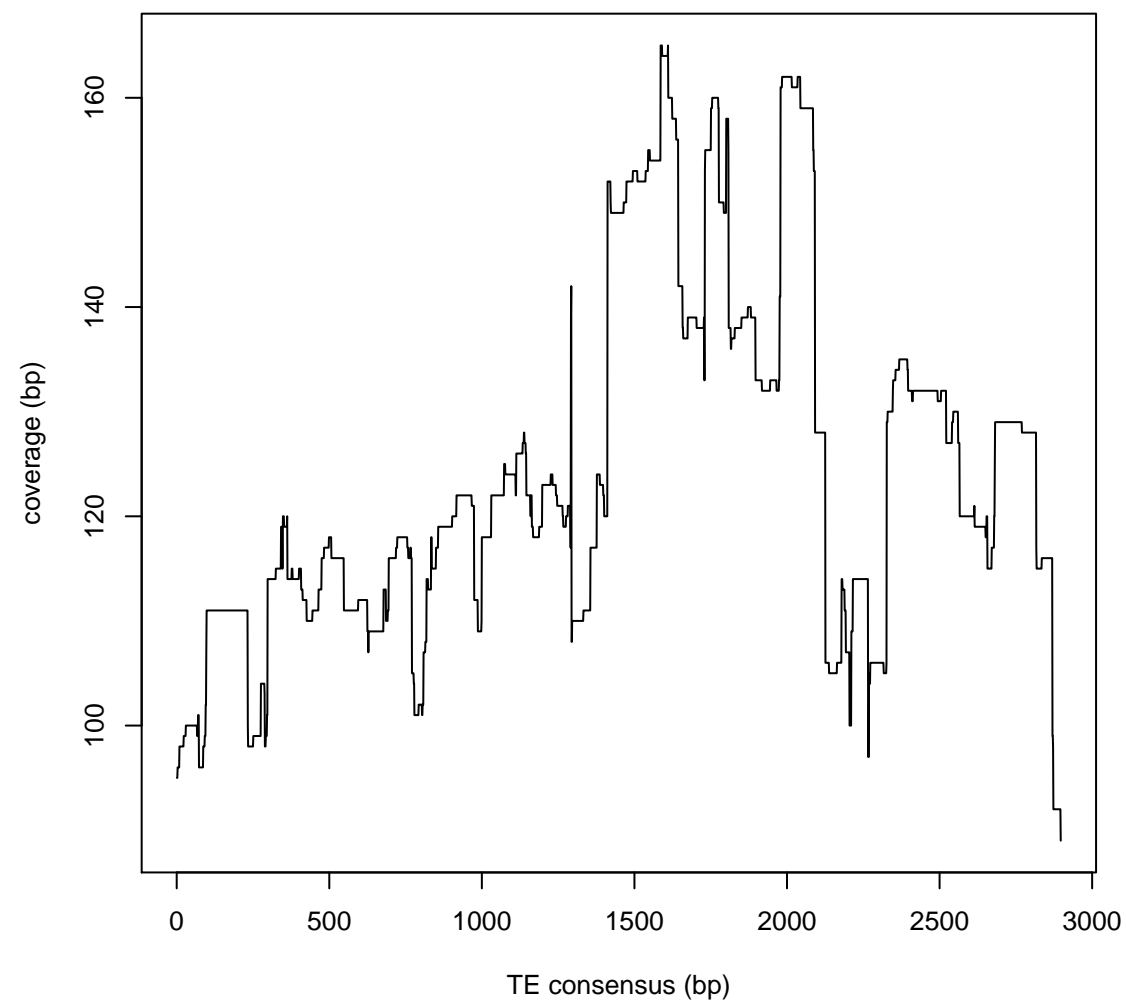


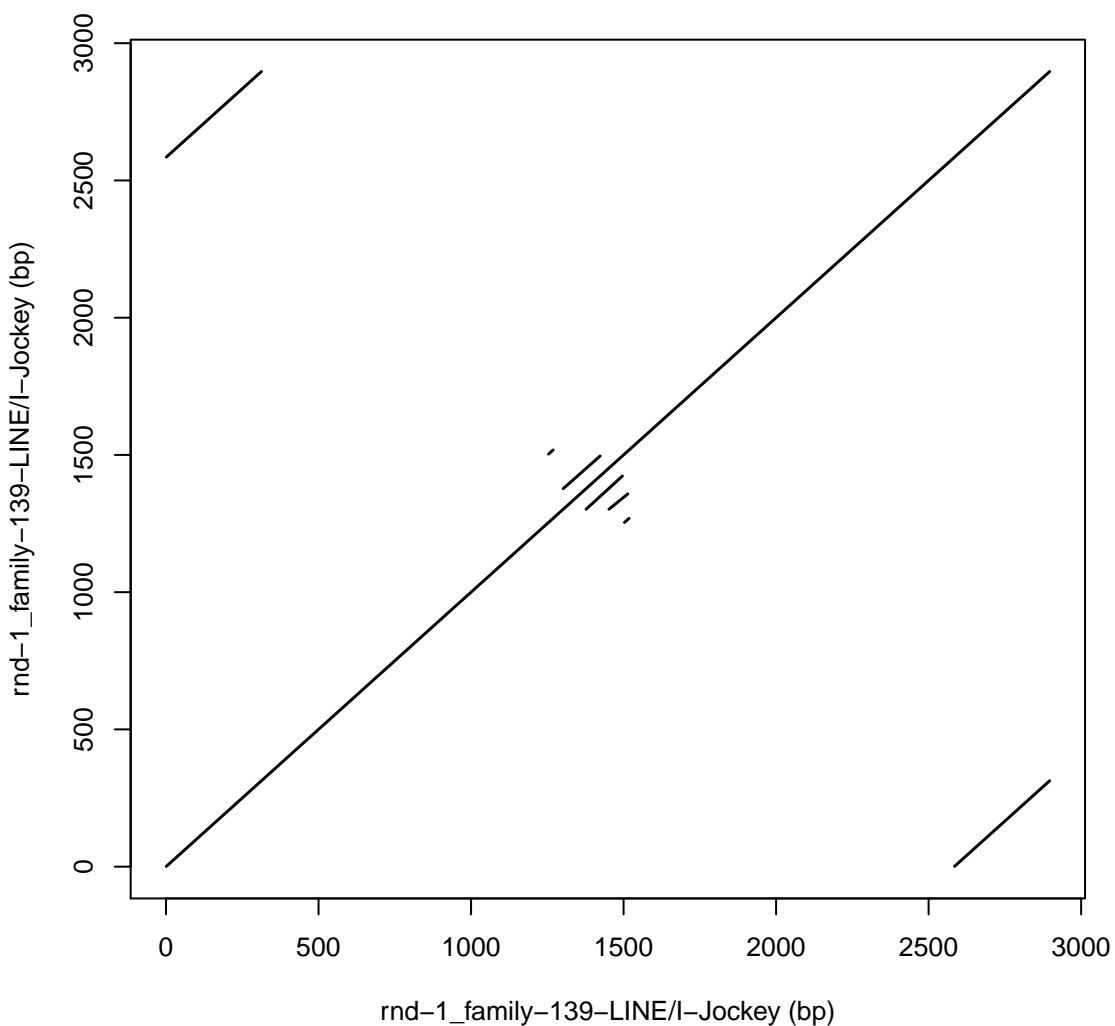
TE: rnd-1_family-139-LINE/I-Jockey
consensus size: 2897bp; fragments: 498; full length: 40 (≥ 2607.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

