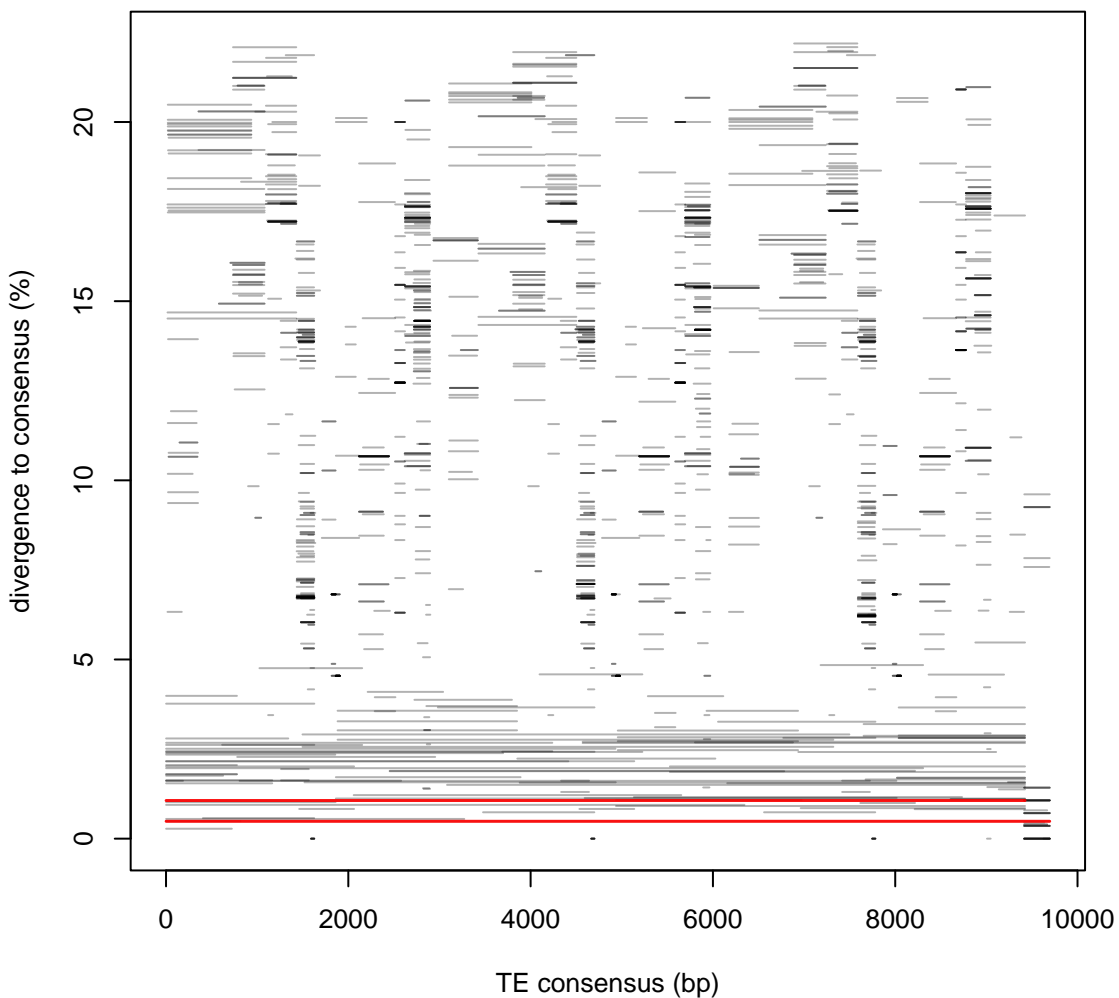
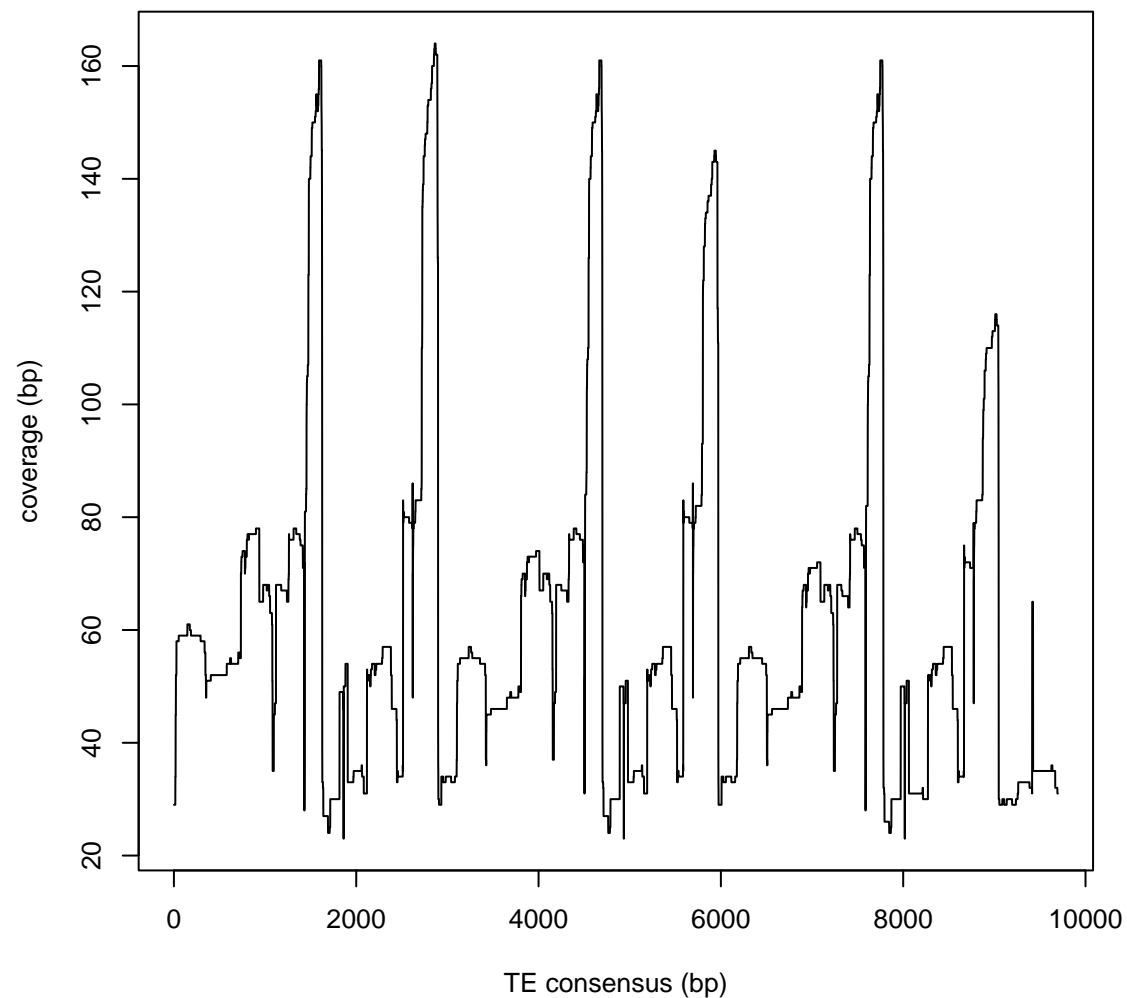


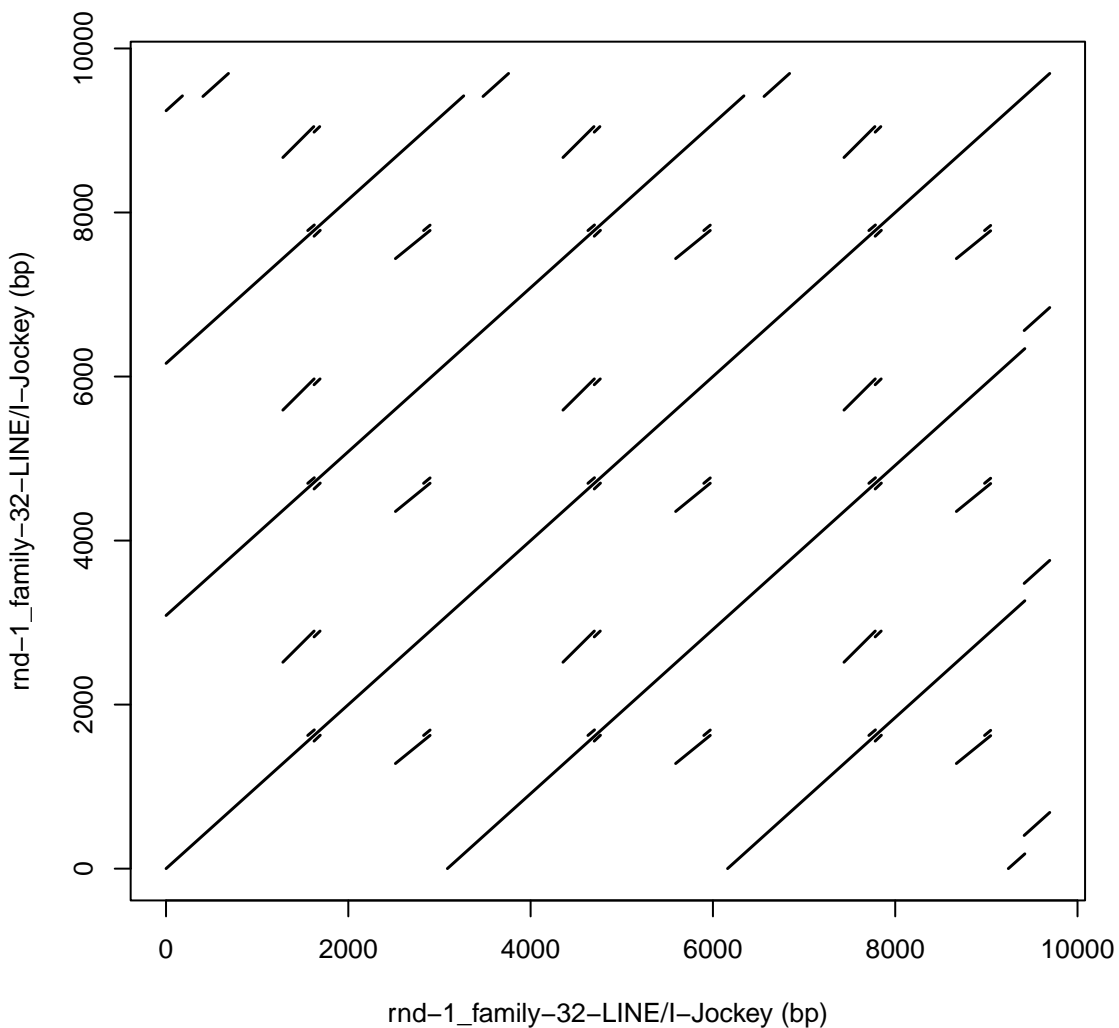
TE: rnd-1_family-32-LINE/I-Jockey
consensus size: 9695bp; fragments: 1622; full length: 2 (≥ 8725.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

