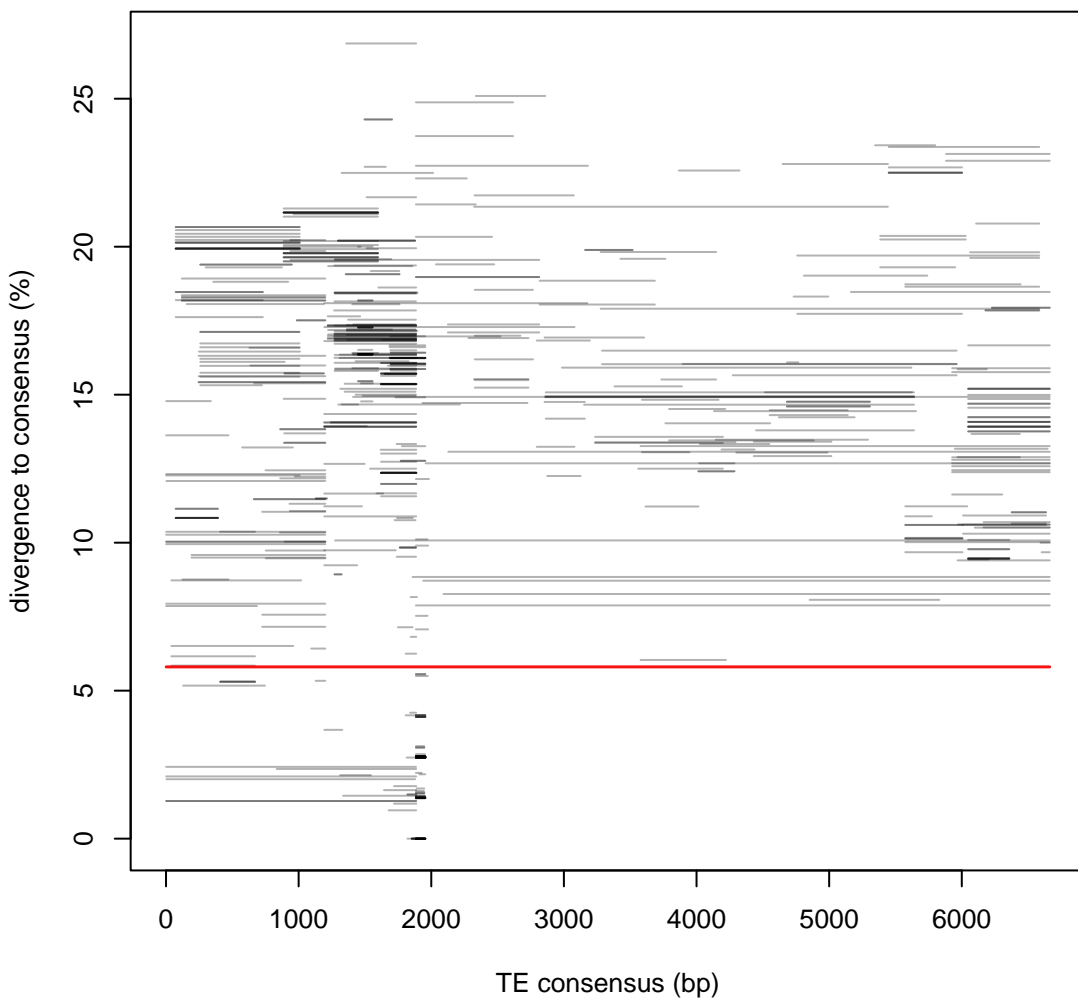
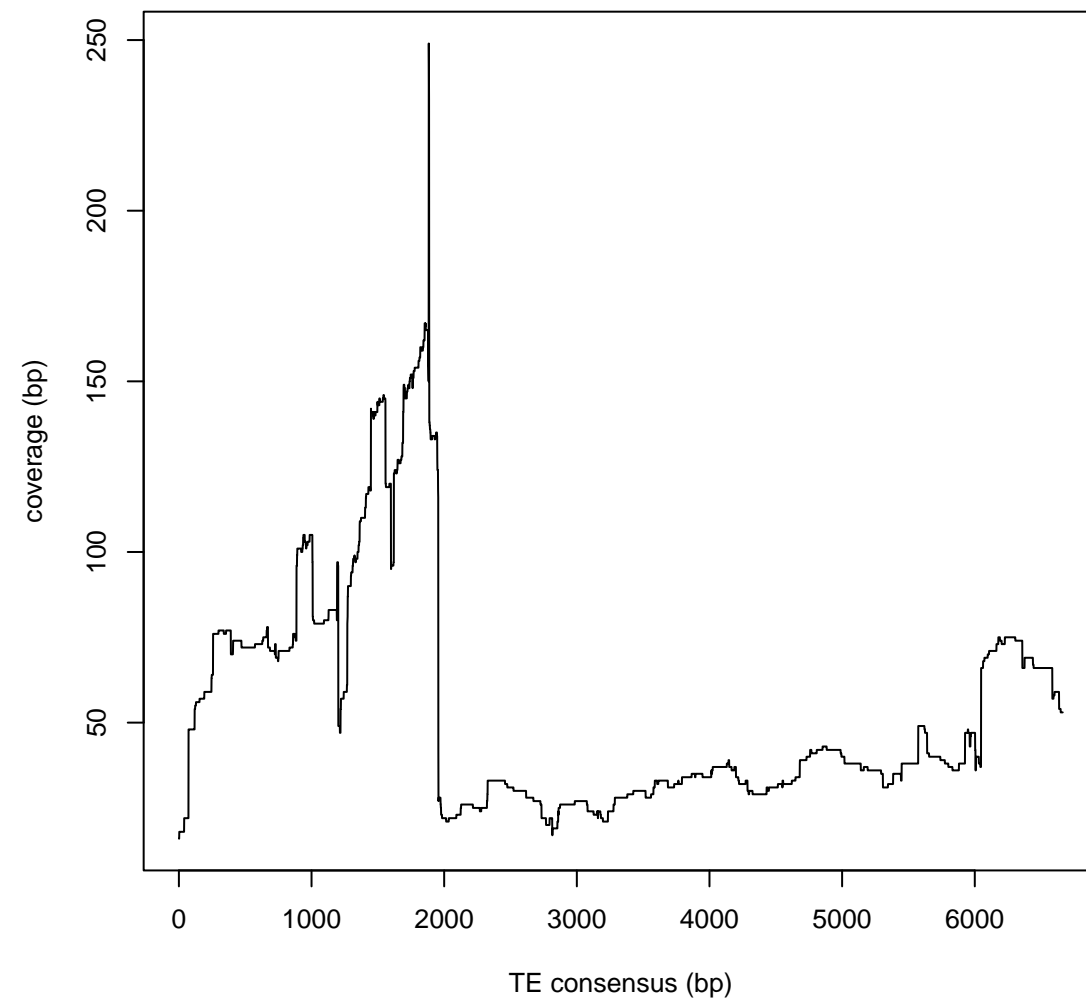


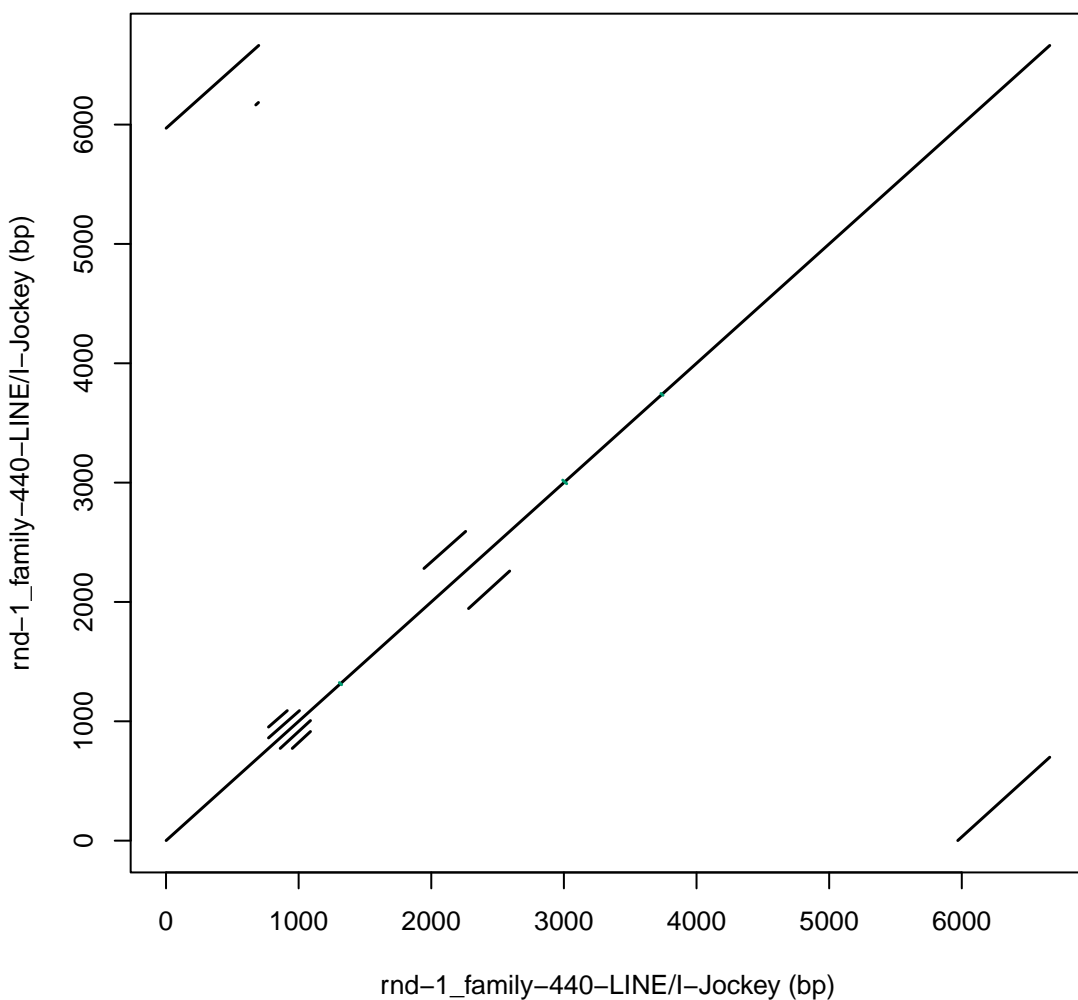
TE: rnd-1_family-440-LINE/I-Jockey
consensus size: 6663bp; fragments: 621; full length: 1 (≥ 5996.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

