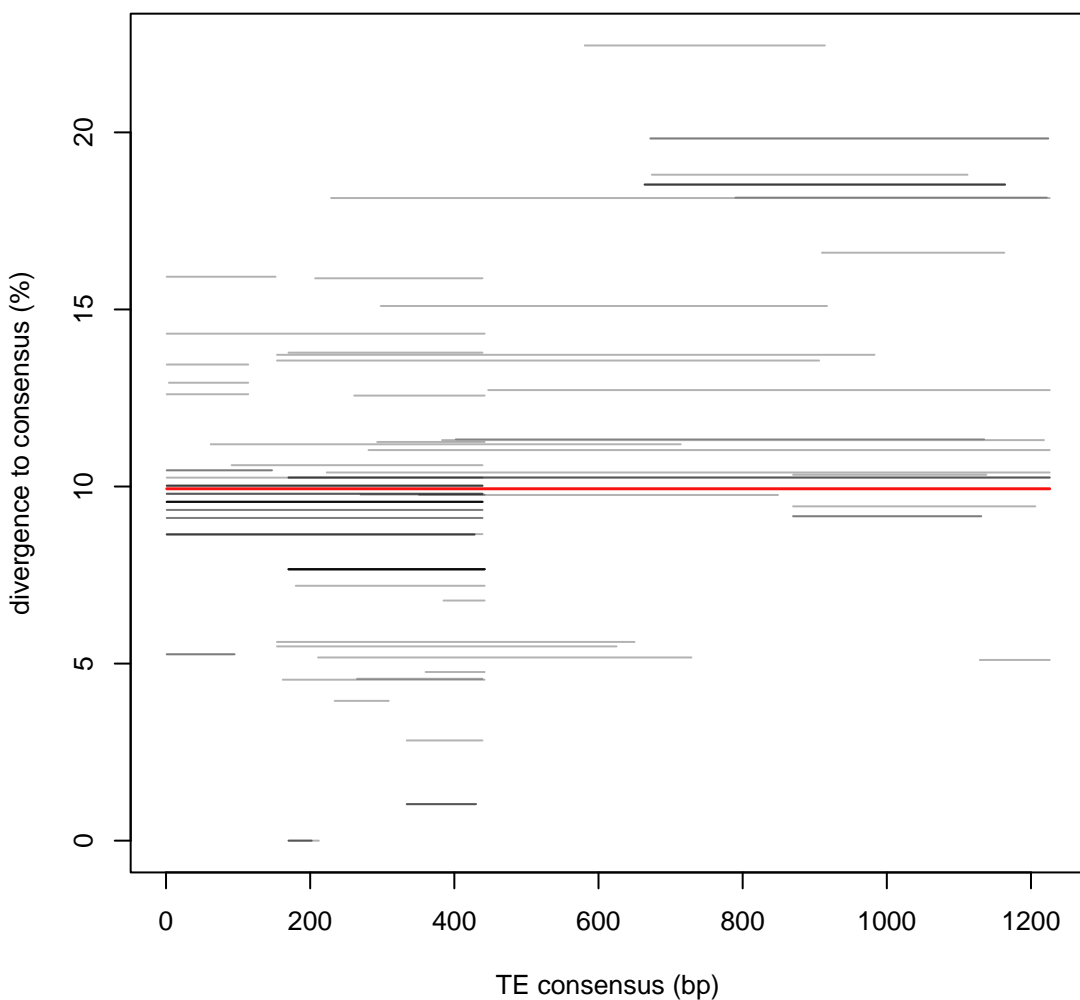
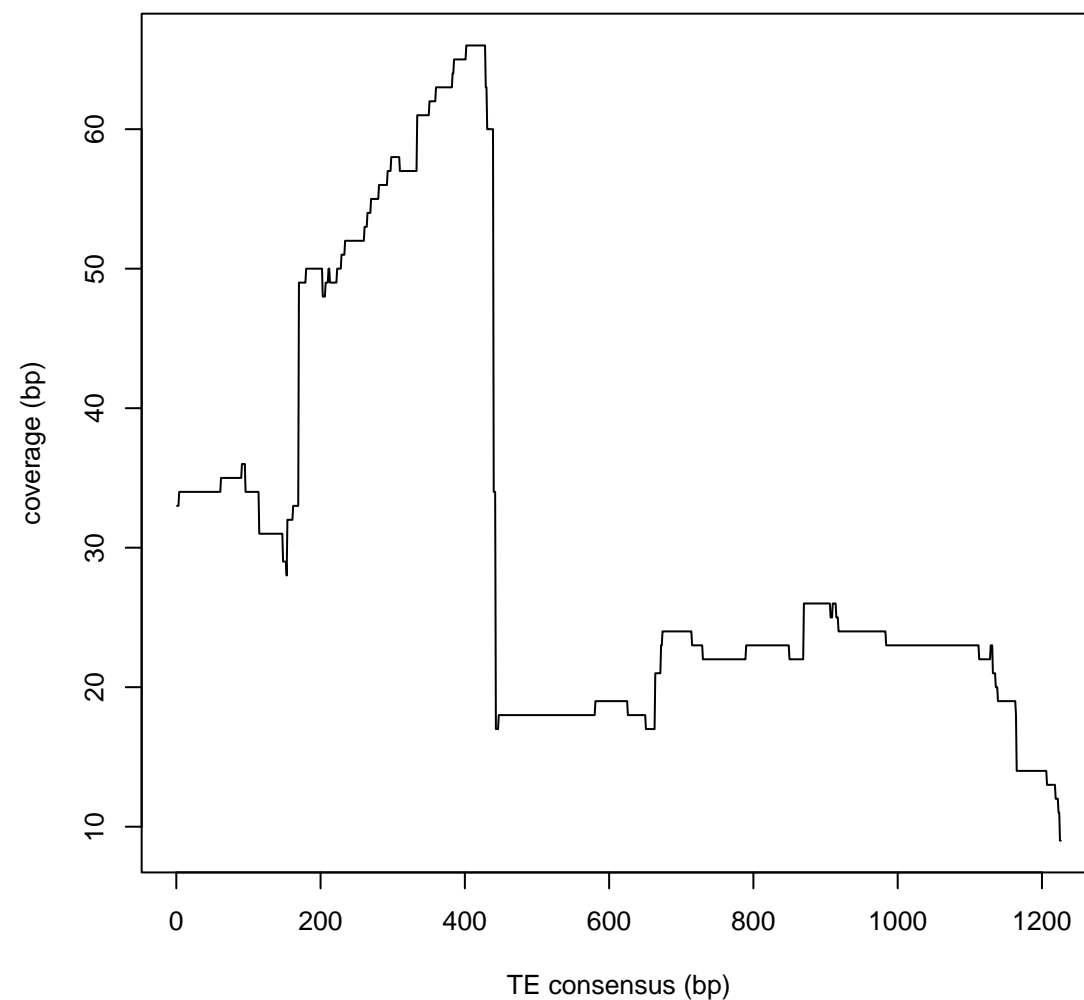


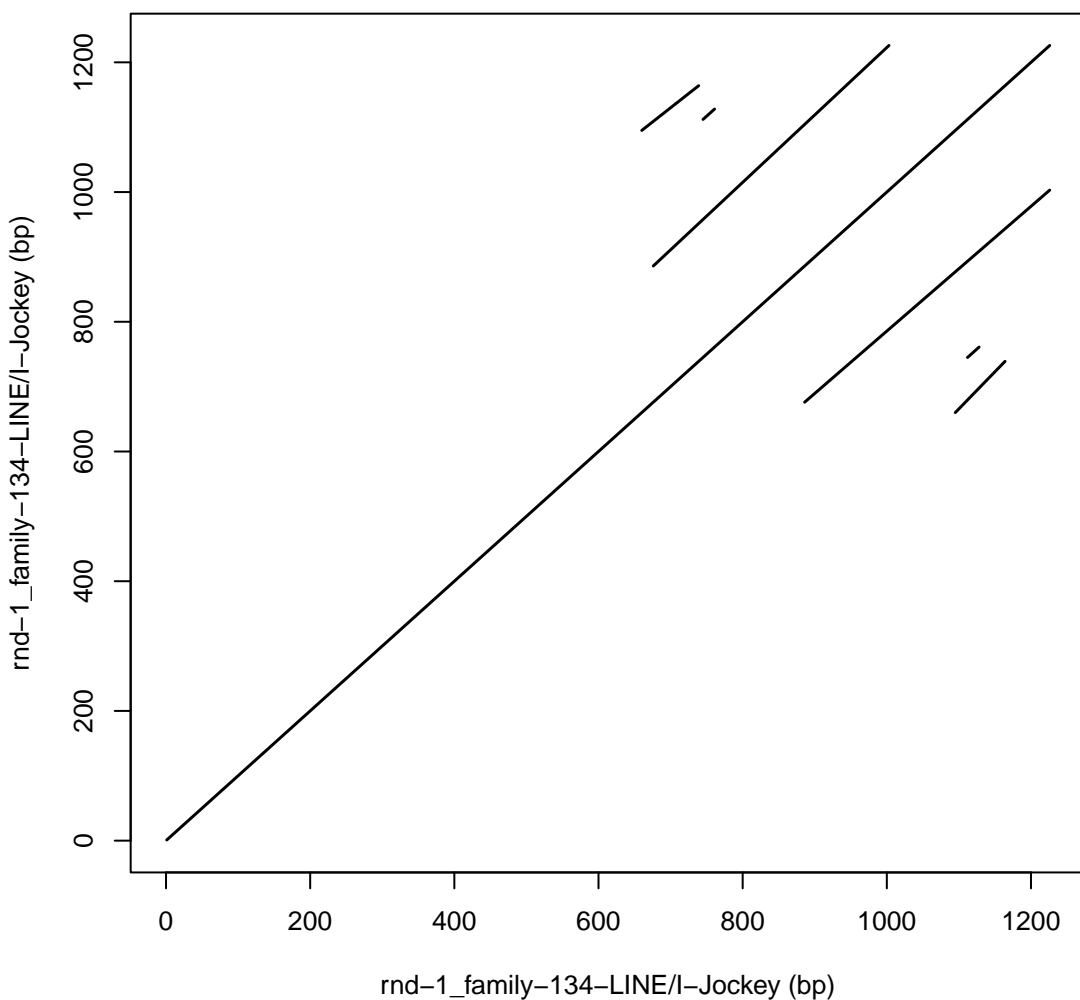
TE: rnd-1_family-134-LINE/I-Jockey
consensus size: 1226bp; fragments: 94; full length: 1 (≥ 1103.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

