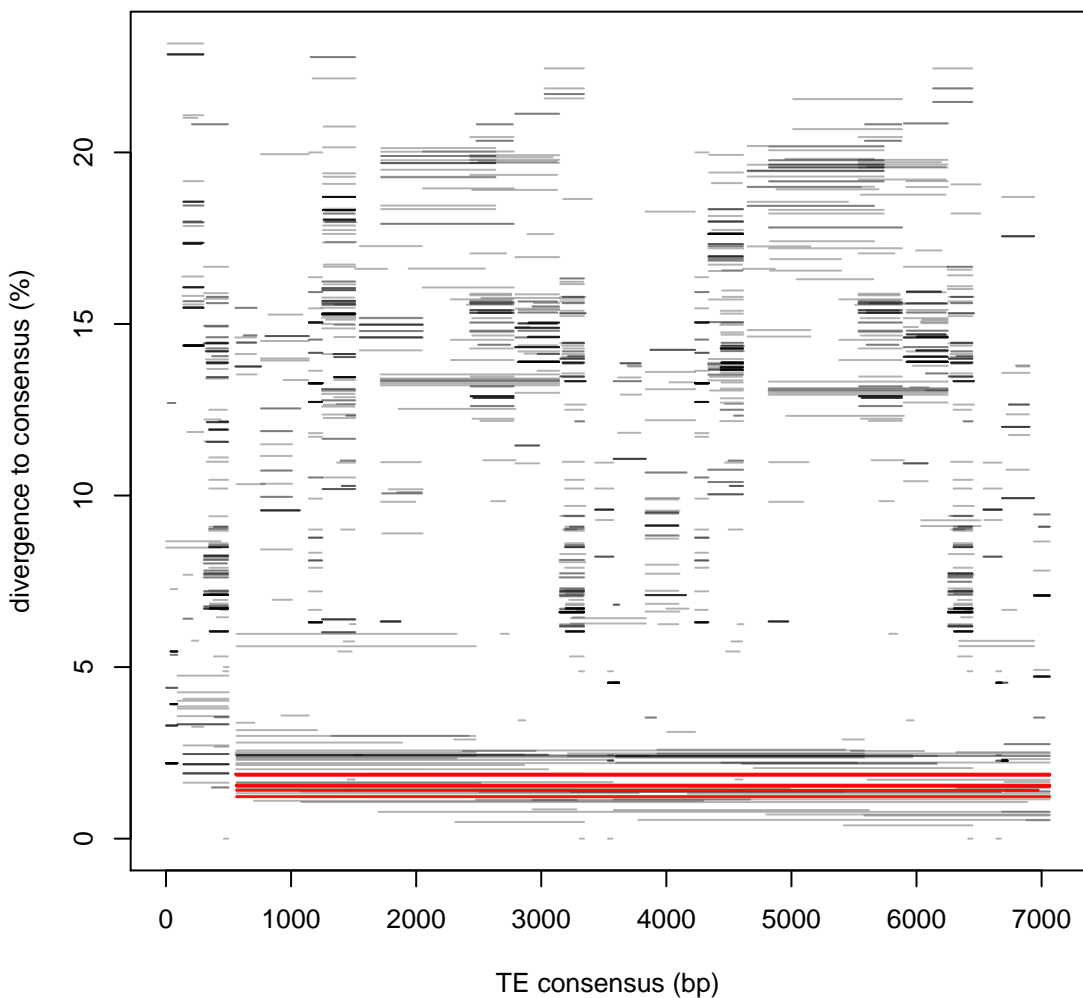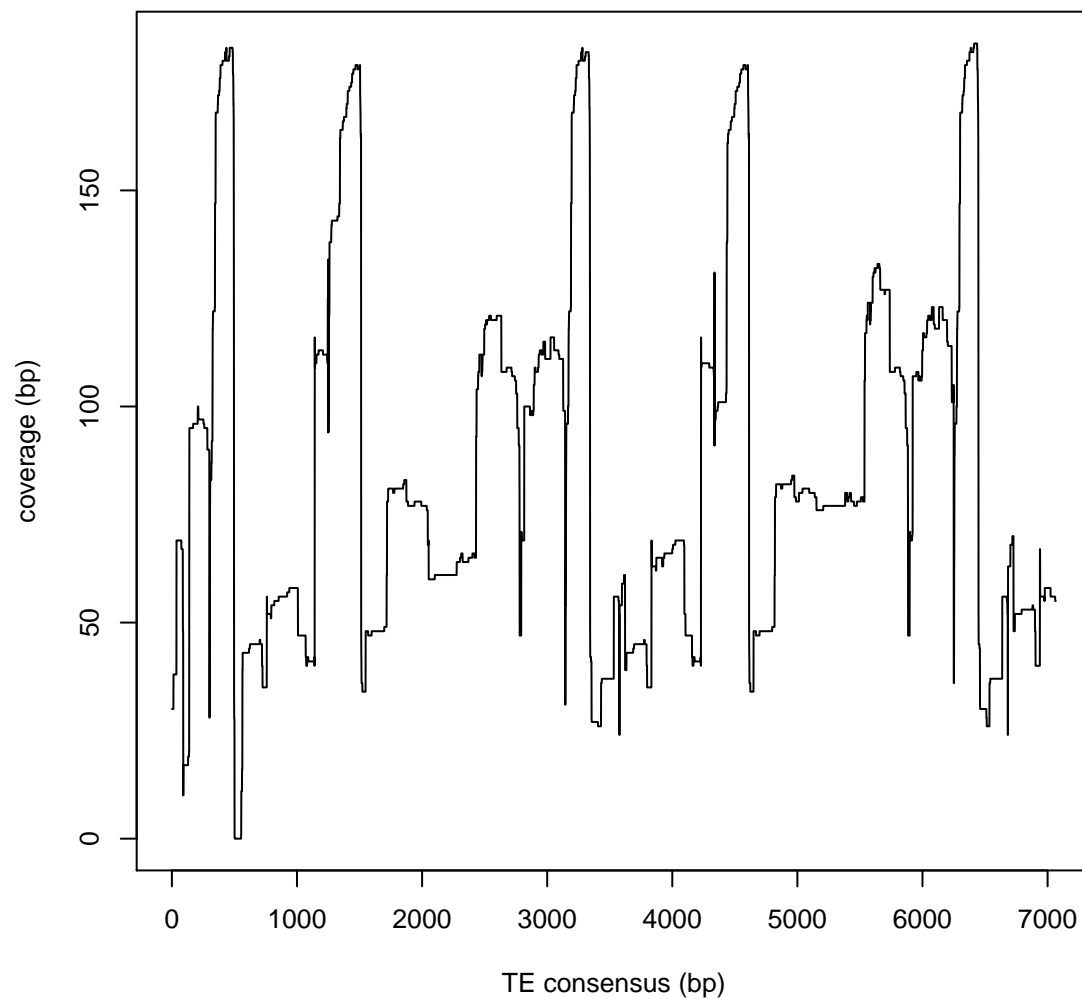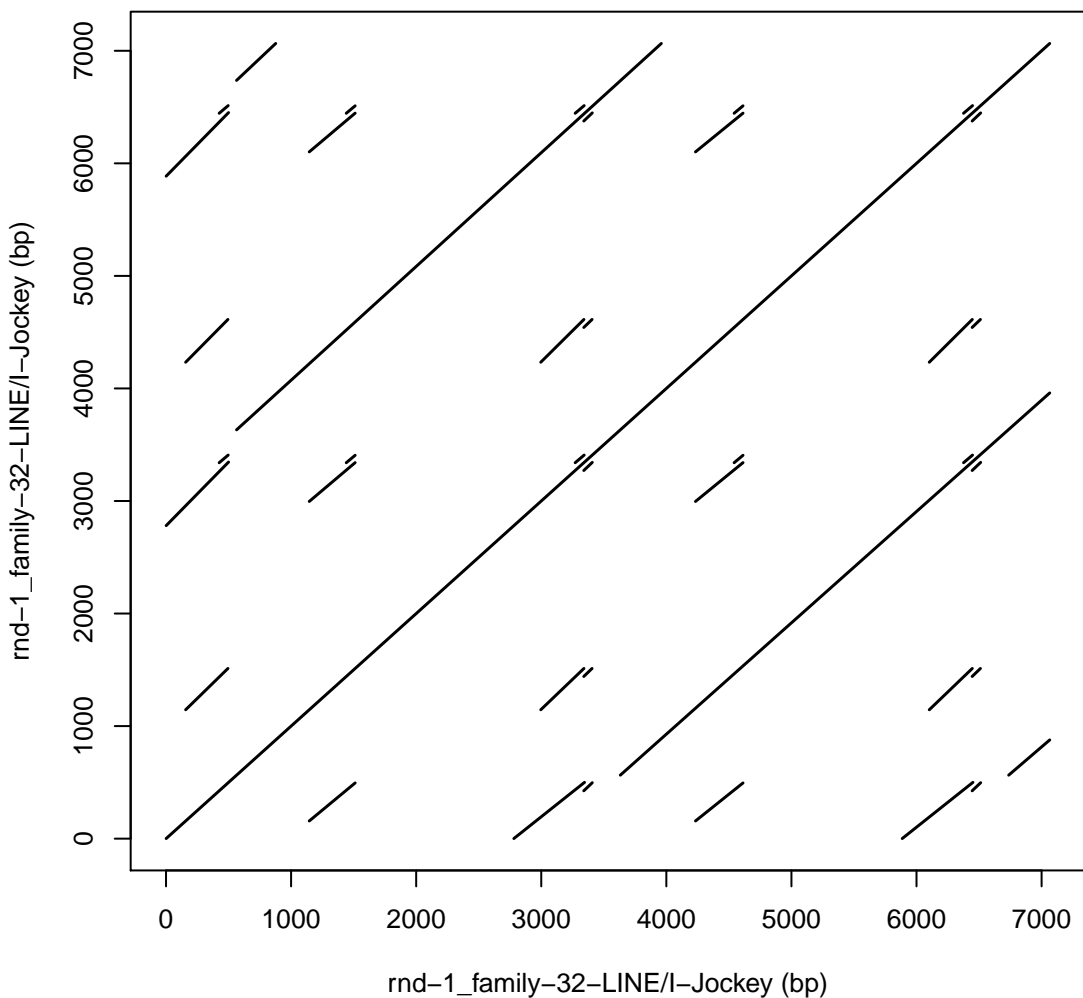# TE: rnd-1_family-32-LINE/I-Jockey
consensus size: 7065bp; fragments: 1813; full length: 6 (>=6358.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected