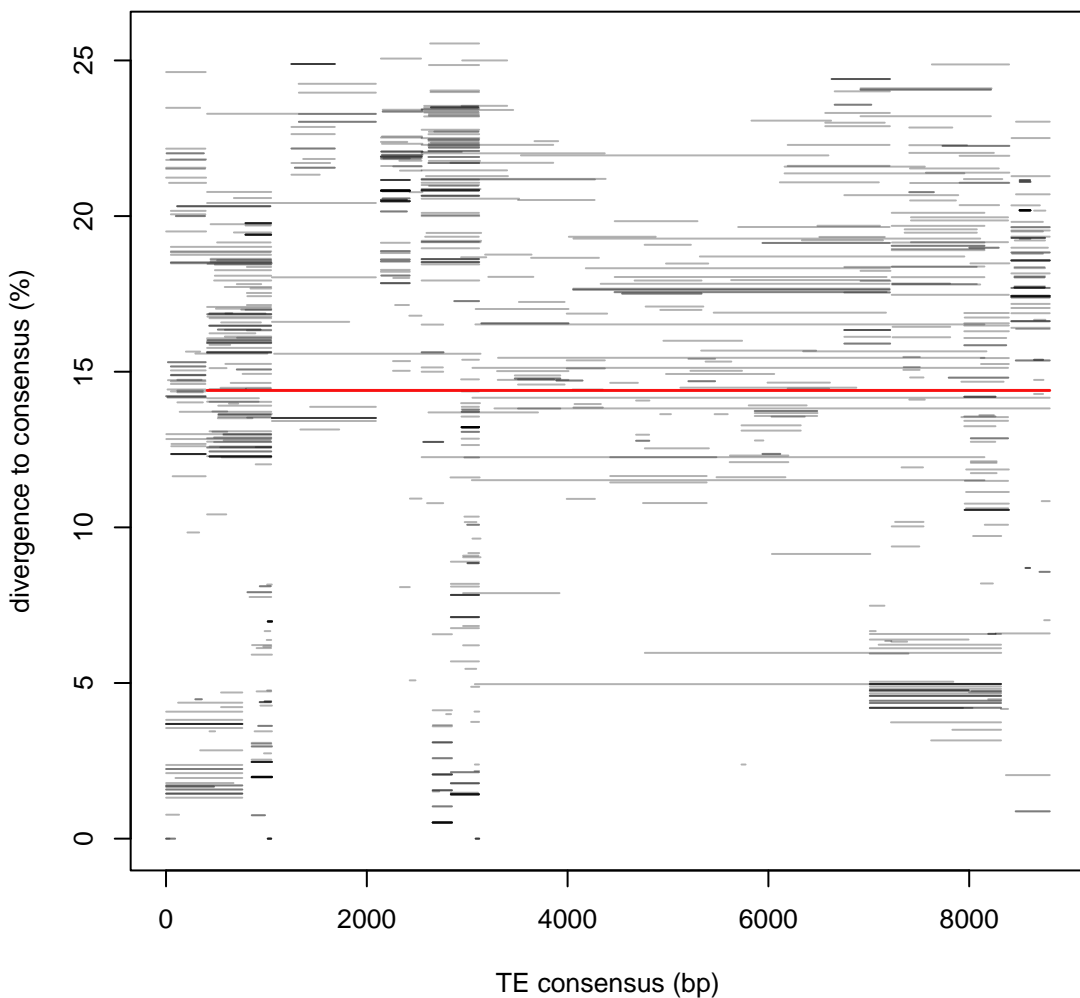
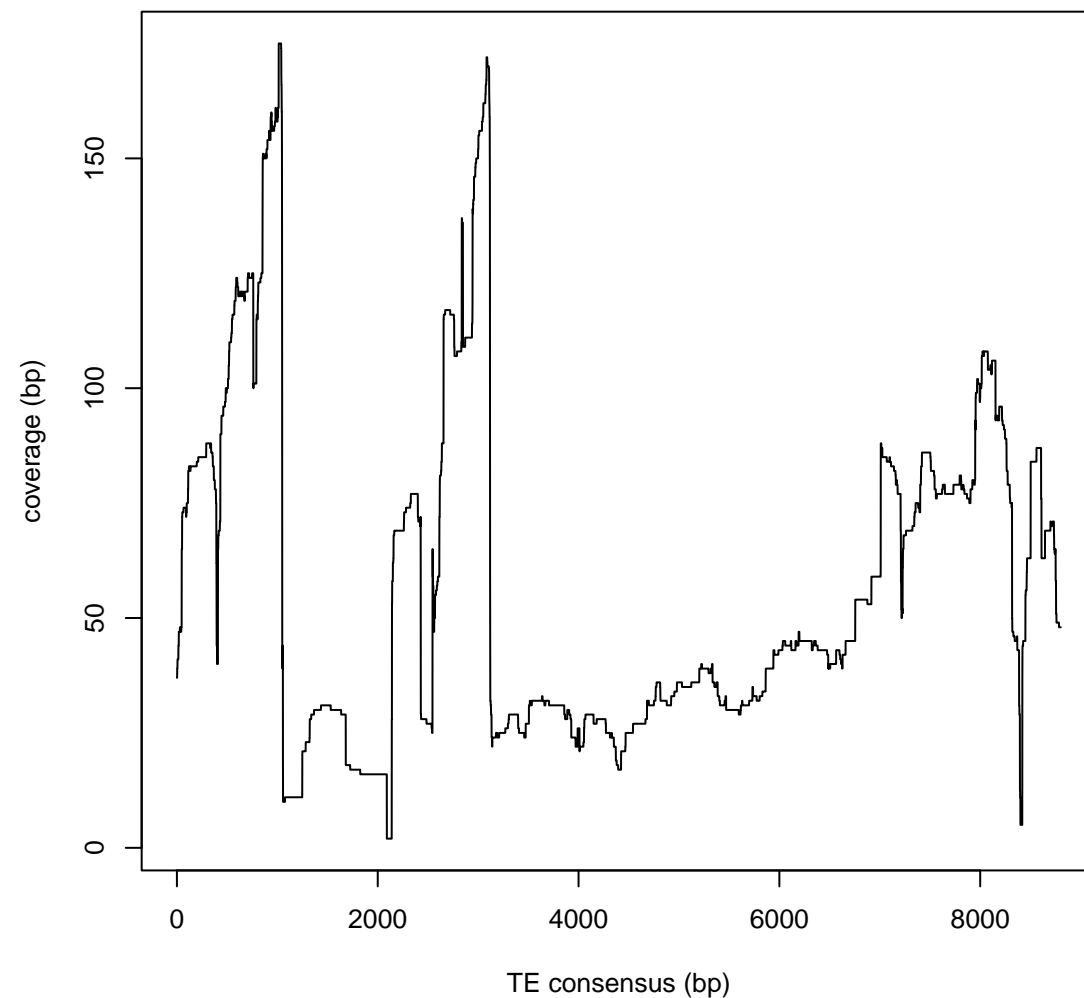


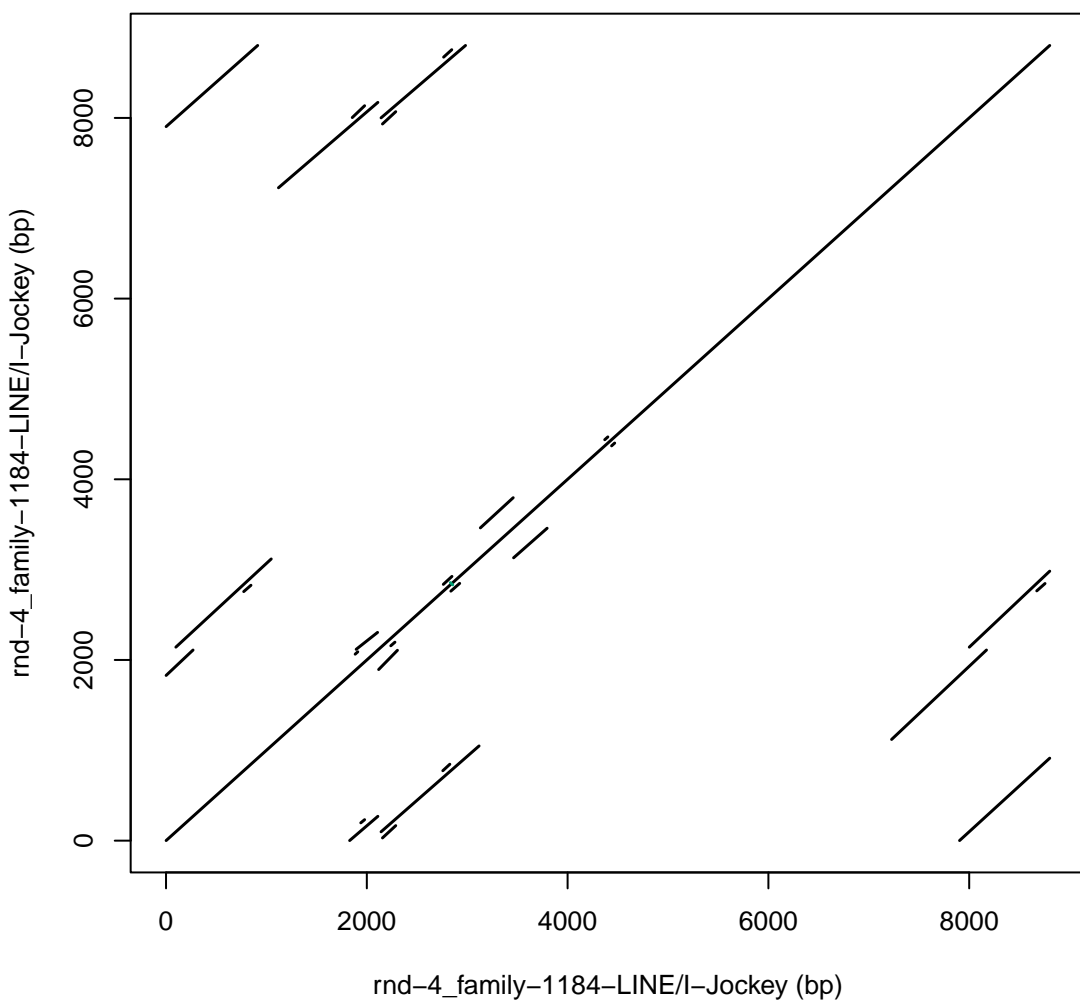
TE: rnd-4_family-1184-LINE/I-Jockey
consensus size: 8802bp; fragments: 942; full length: 1 (>=7921.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

