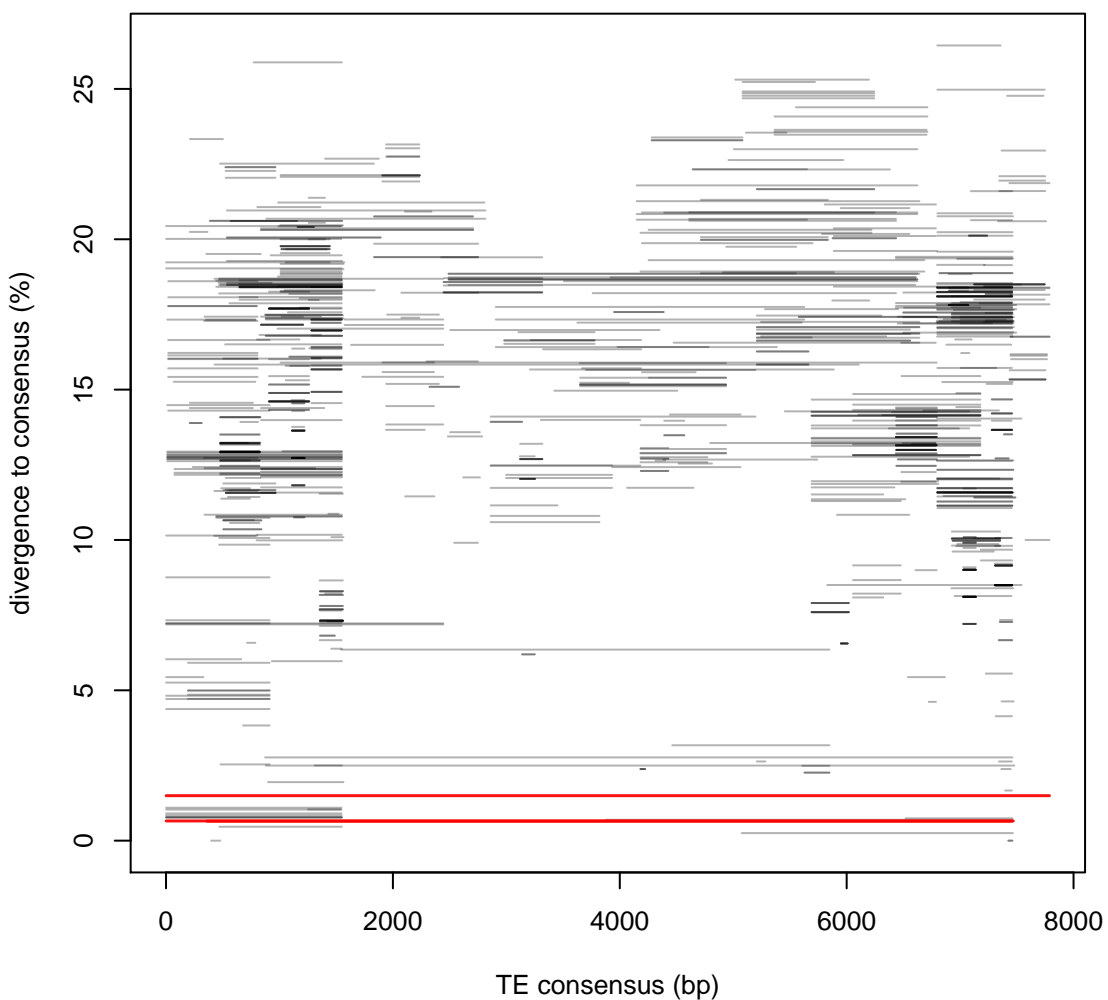
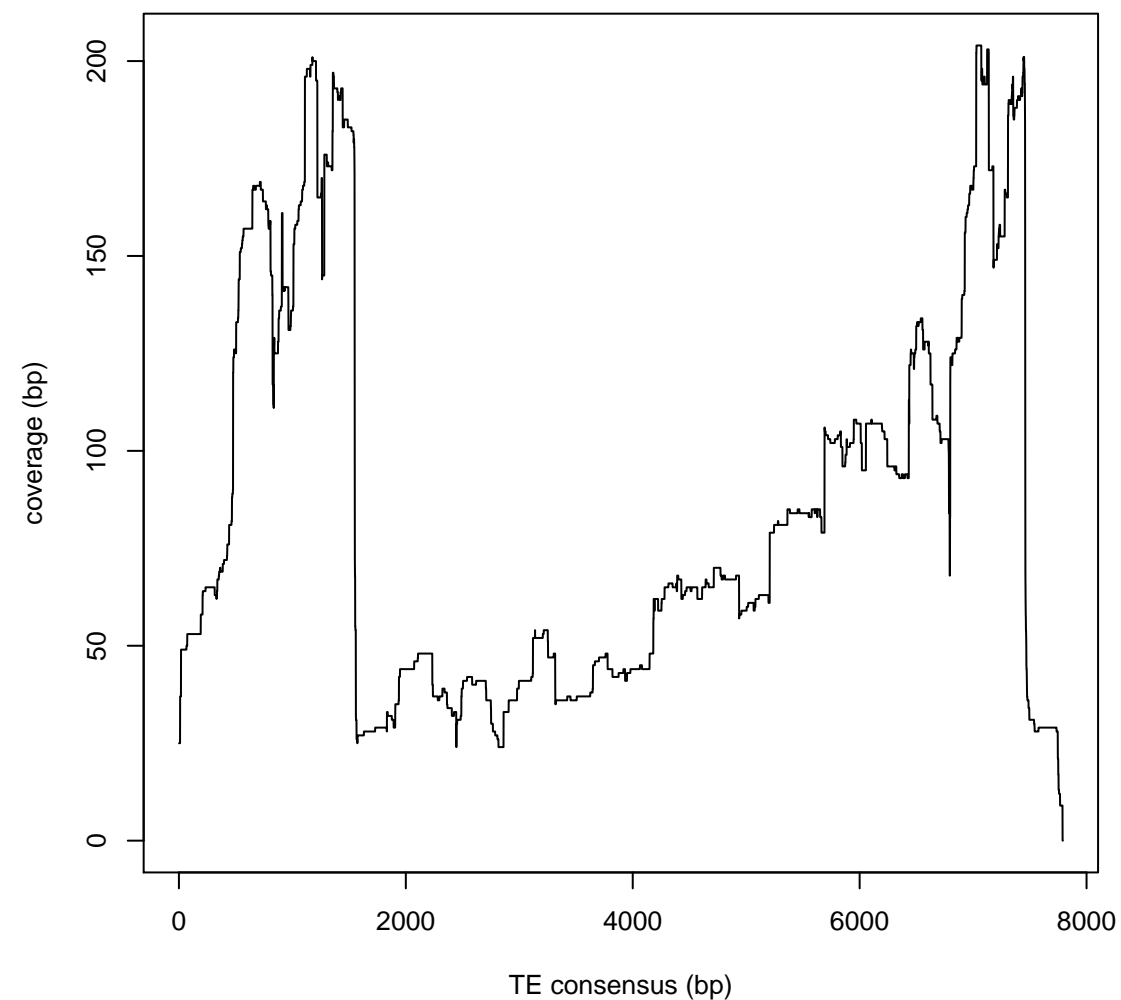


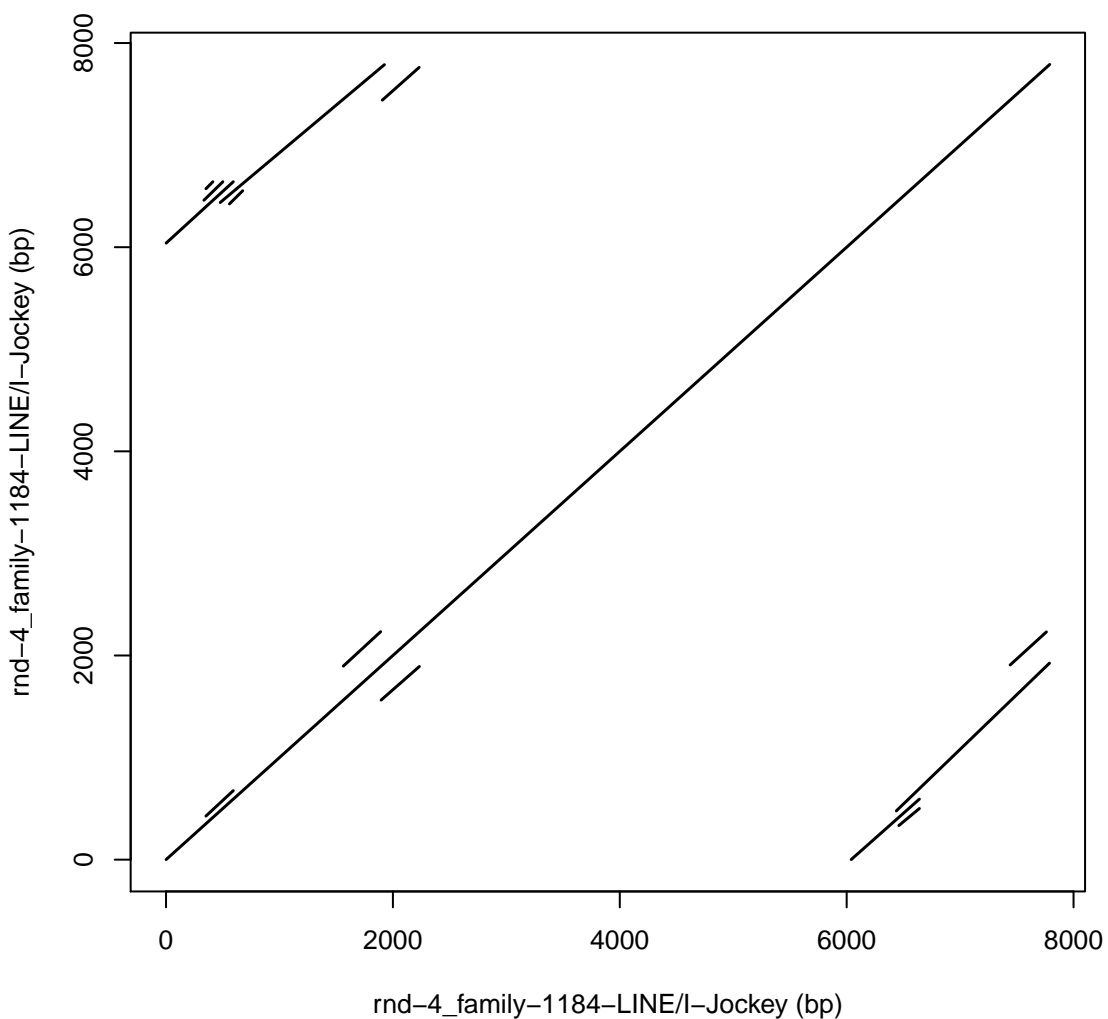
TE: rnd-4_family-1184-LINE/I-Jockey
consensus size: 7790bp; fragments: 947; full length: 3 (>=7011bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

