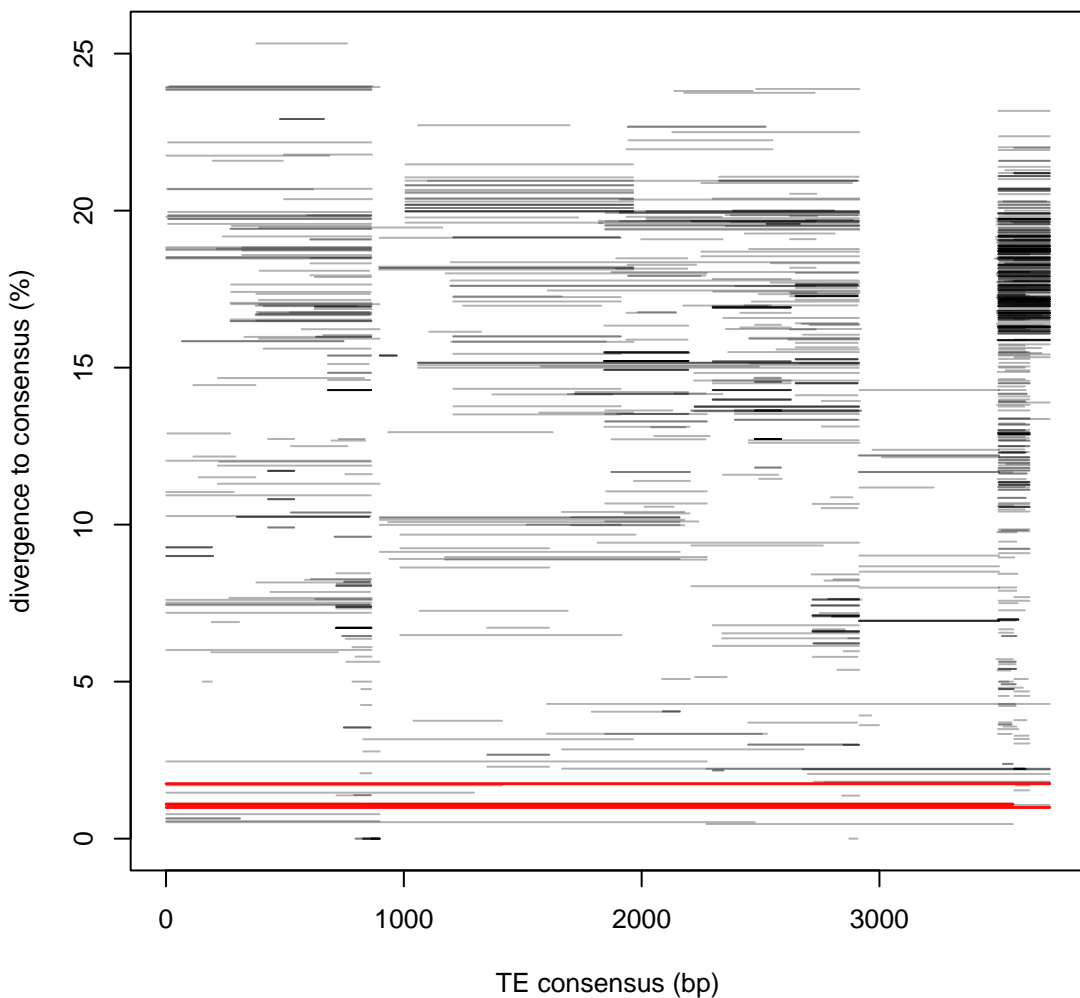
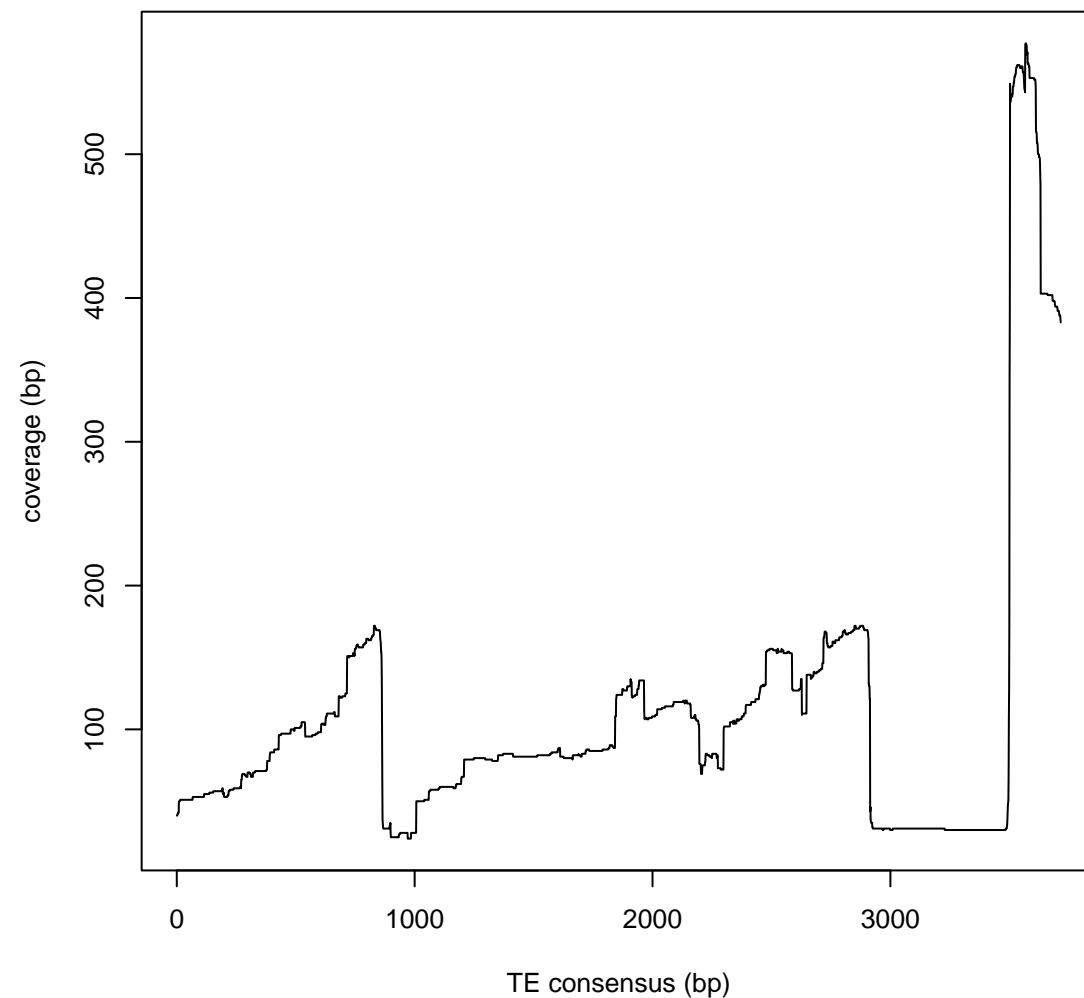


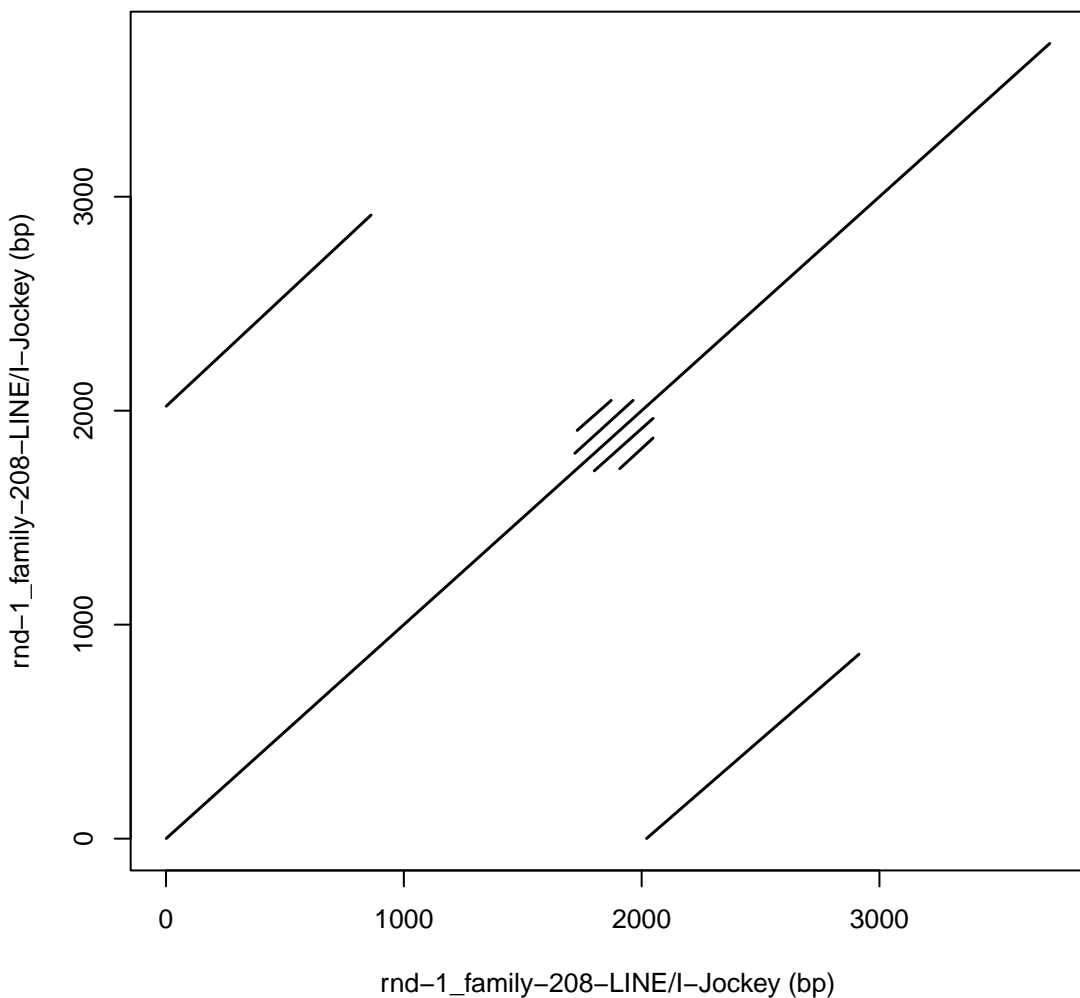
TE: rnd-1_family-208-LINE/I-Jockey
consensus size: 3716bp; fragments: 1215; full length: 4 (>=3344.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

