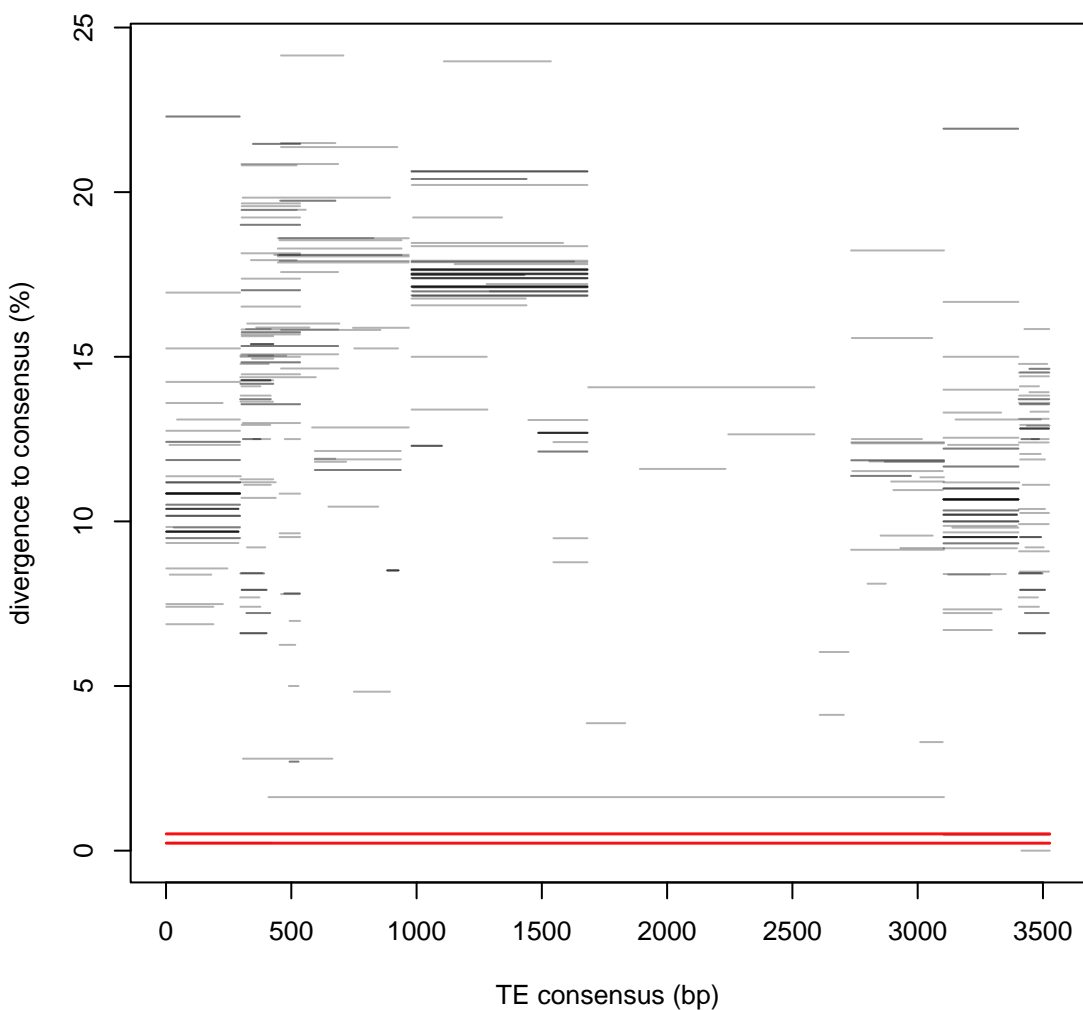
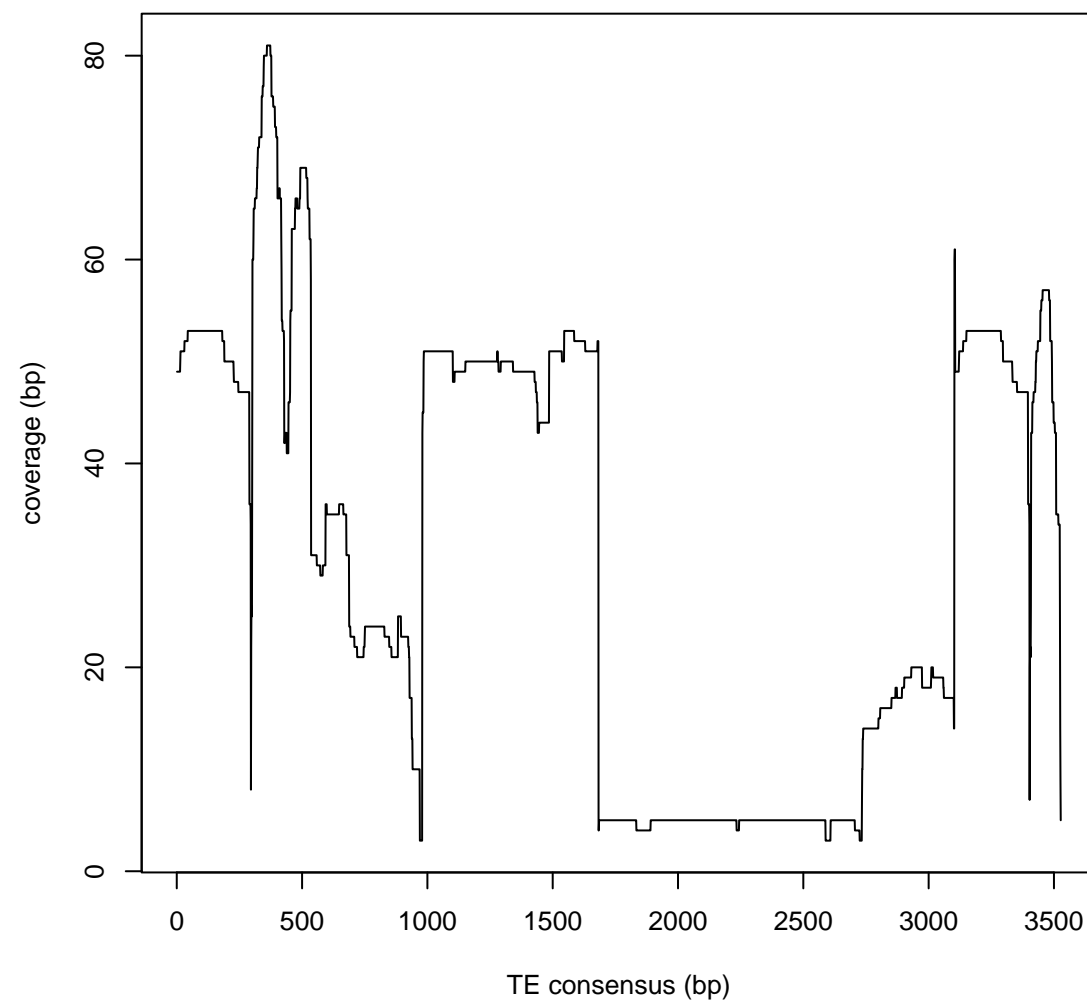


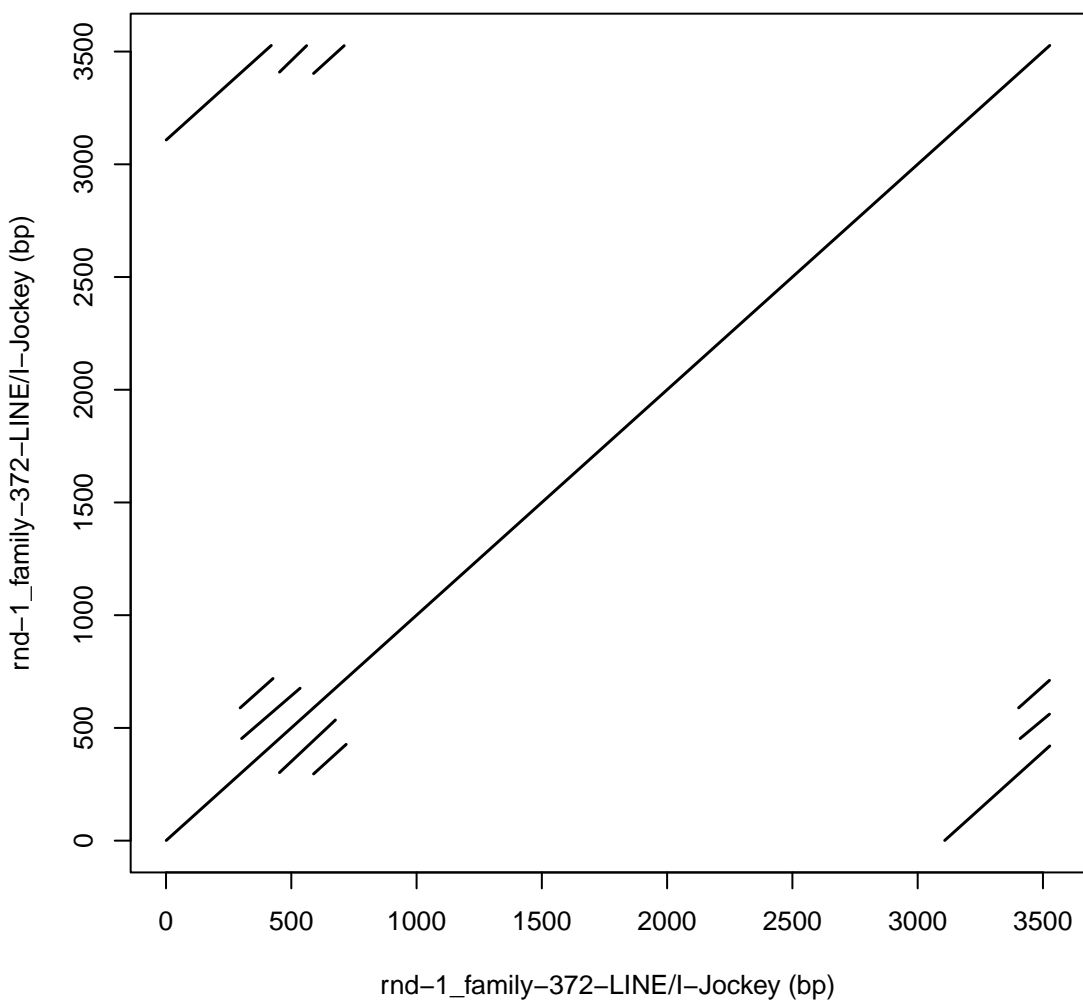
TE: rnd-1_family-372-LINE/I-Jockey
consensus size: 3527bp; fragments: 371; full length: 2 (≥ 3174.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

