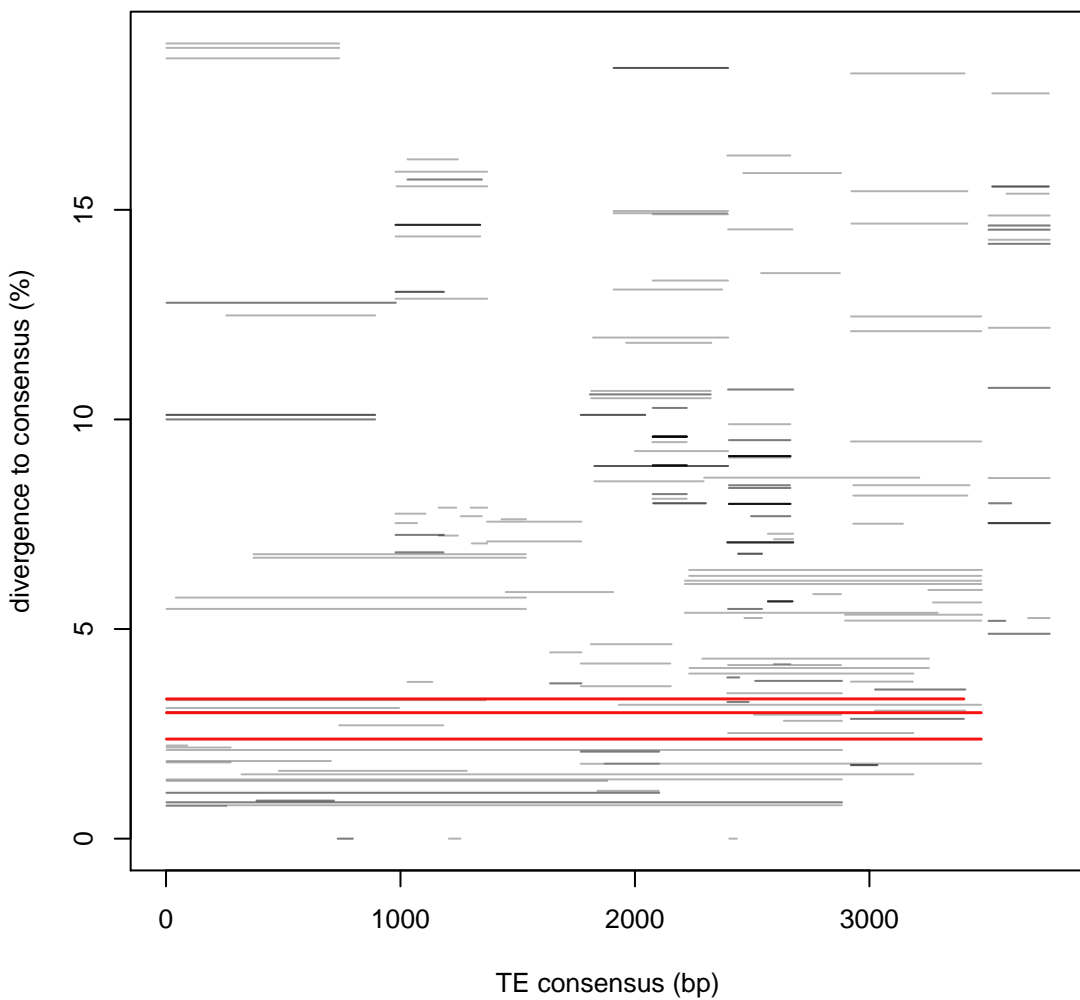
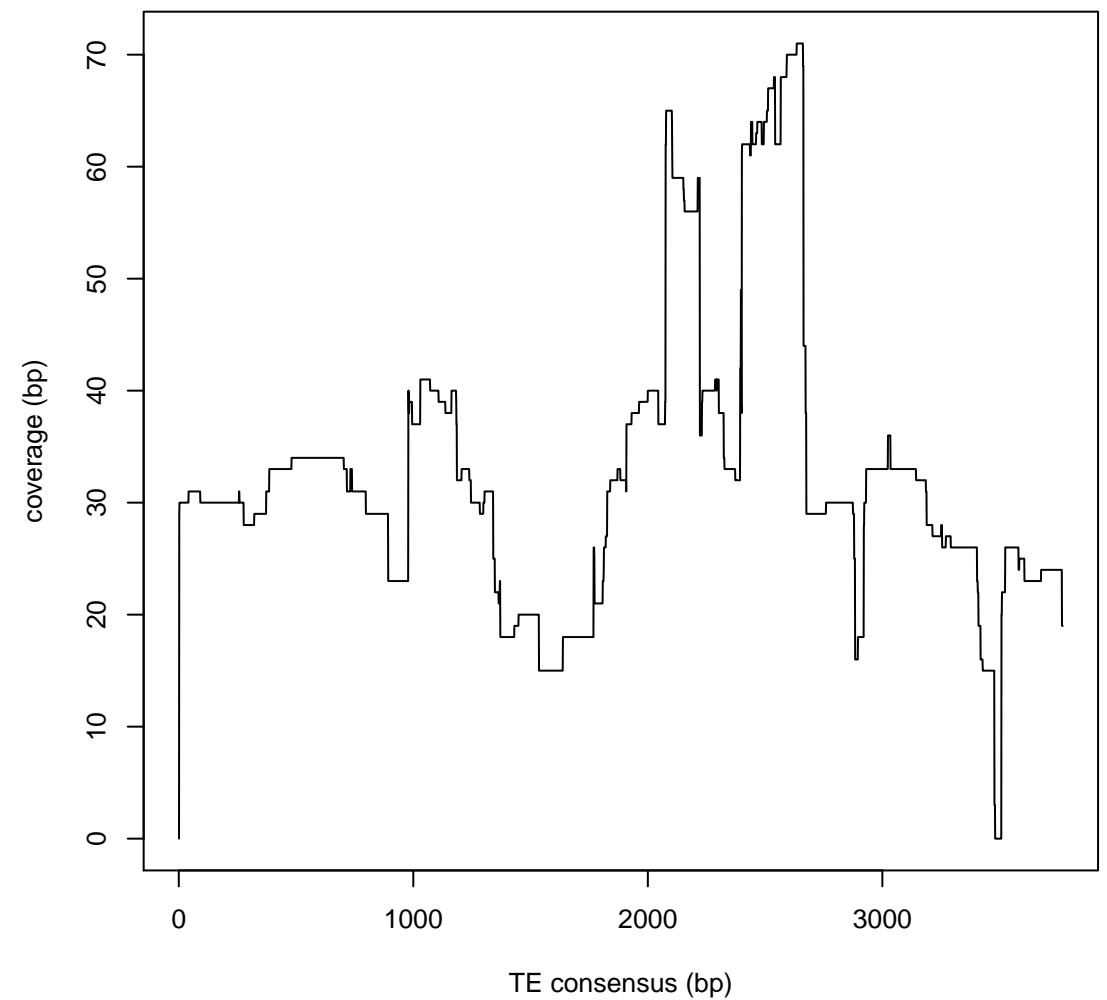


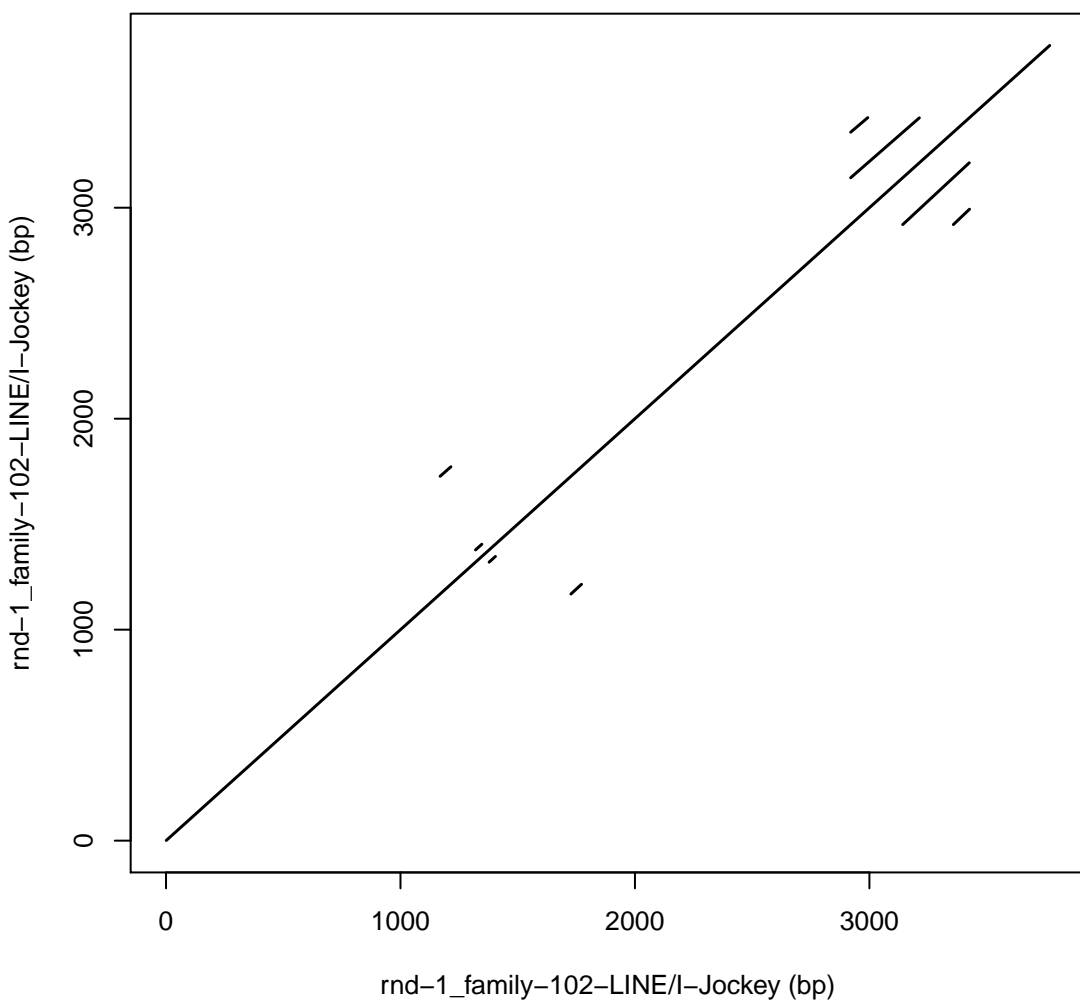
TE: rnd-1_family-102-LINE/I-Jockey
consensus size: 3769bp; fragments: 255; full length: 3 (≥ 3392.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

