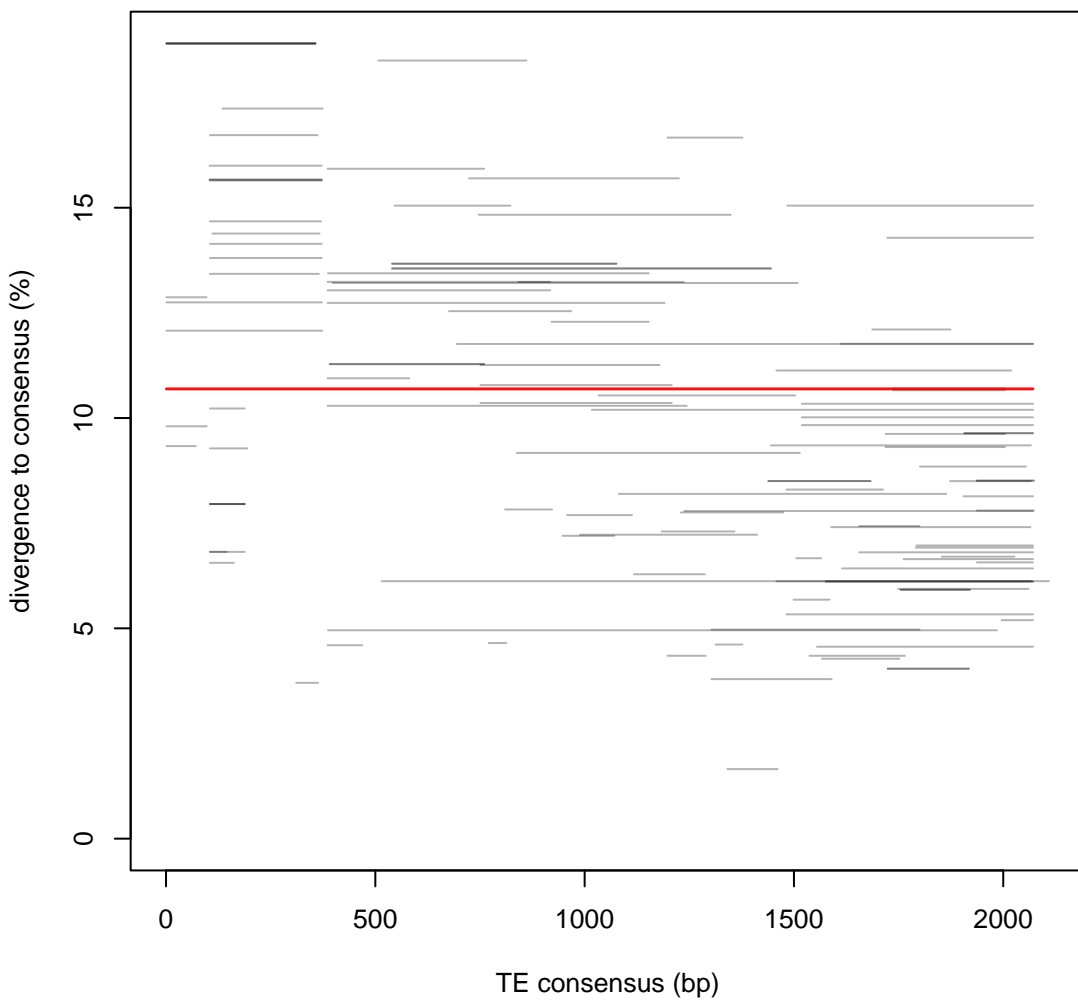
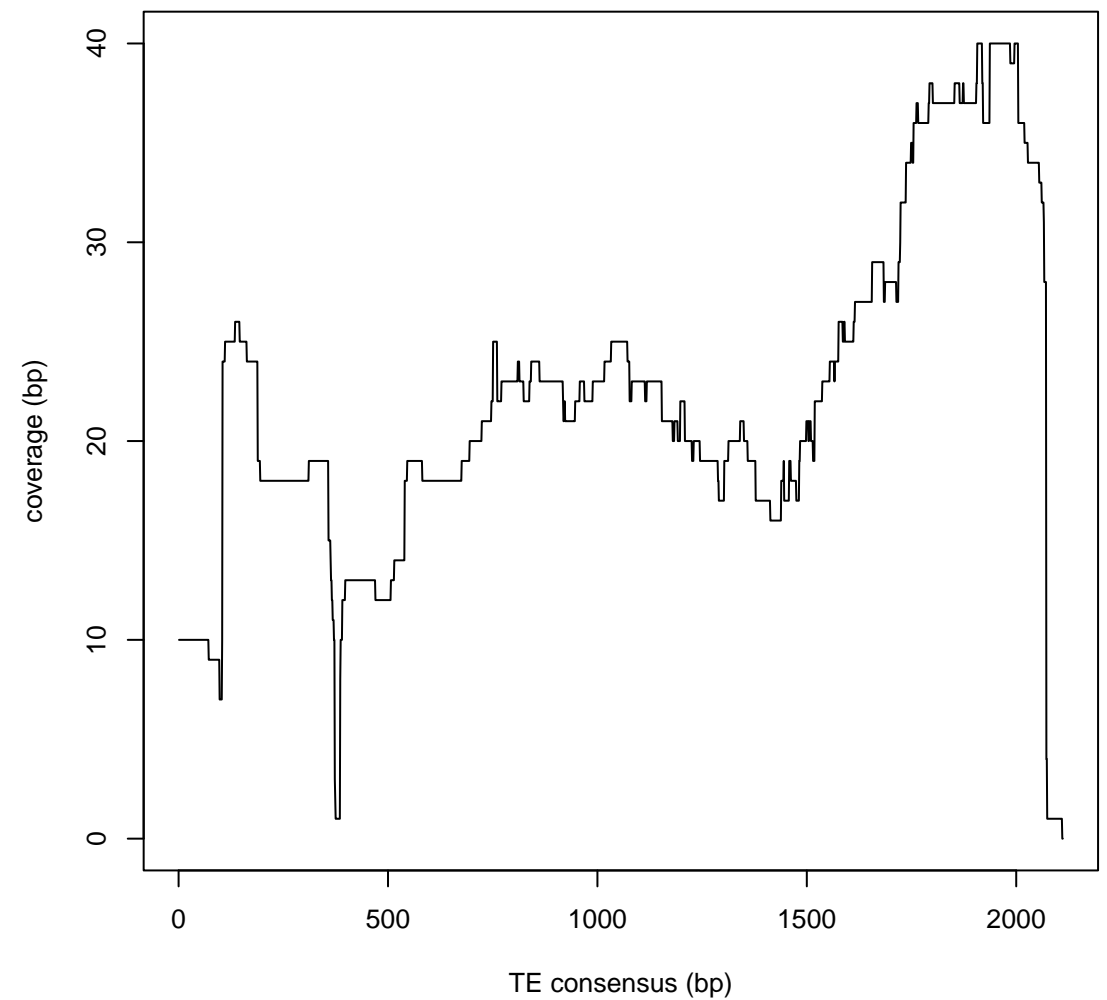


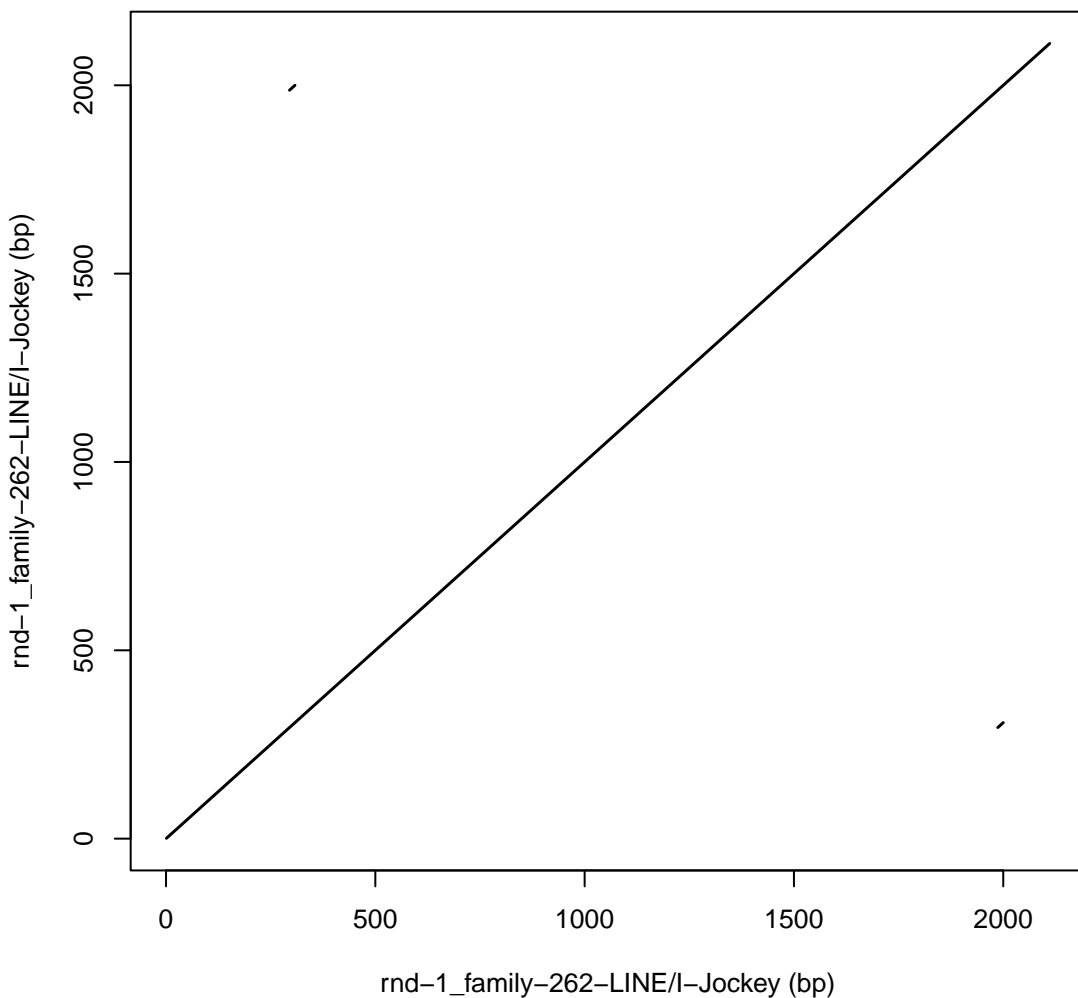
TE: rnd-1_family-262-LINE/I-Jockey
consensus size: 2111bp; fragments: 125; full length: 1 (≥ 1899.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

