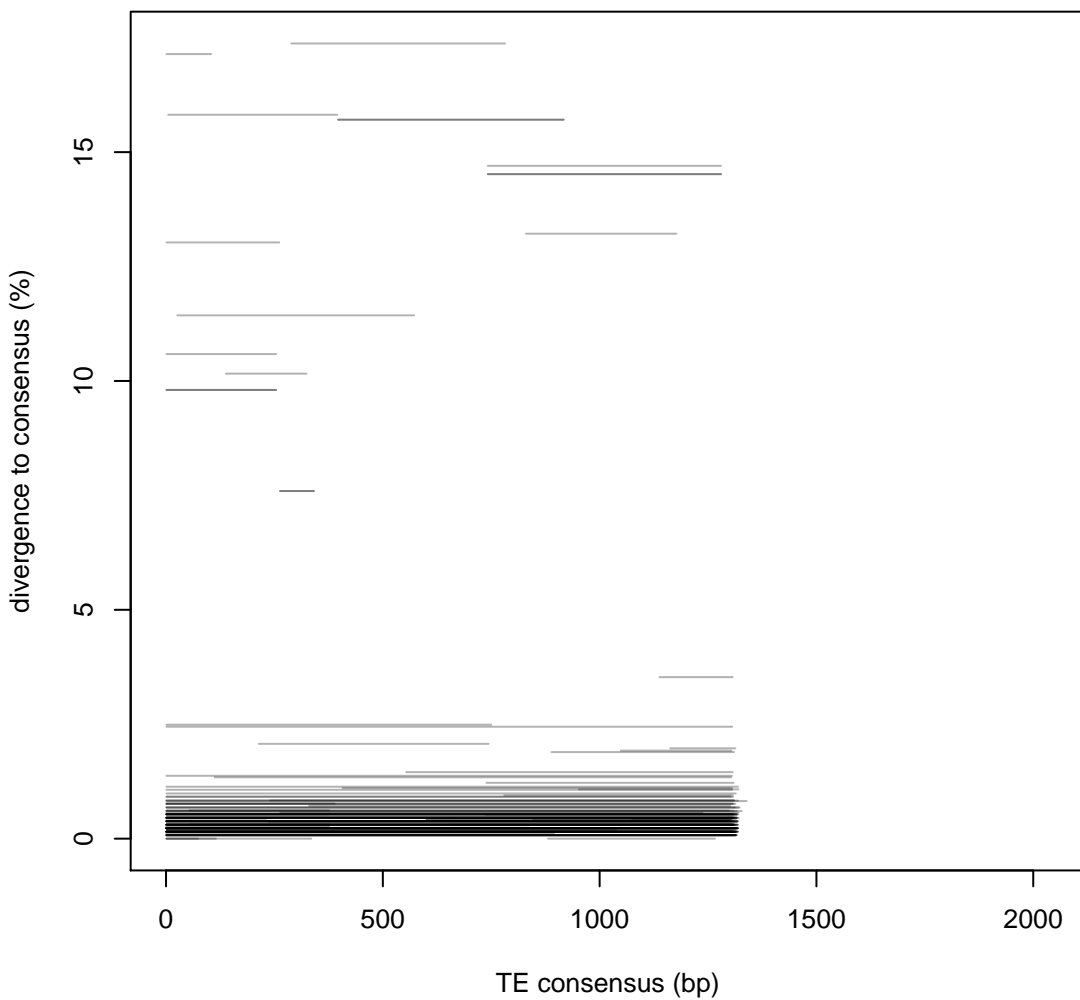
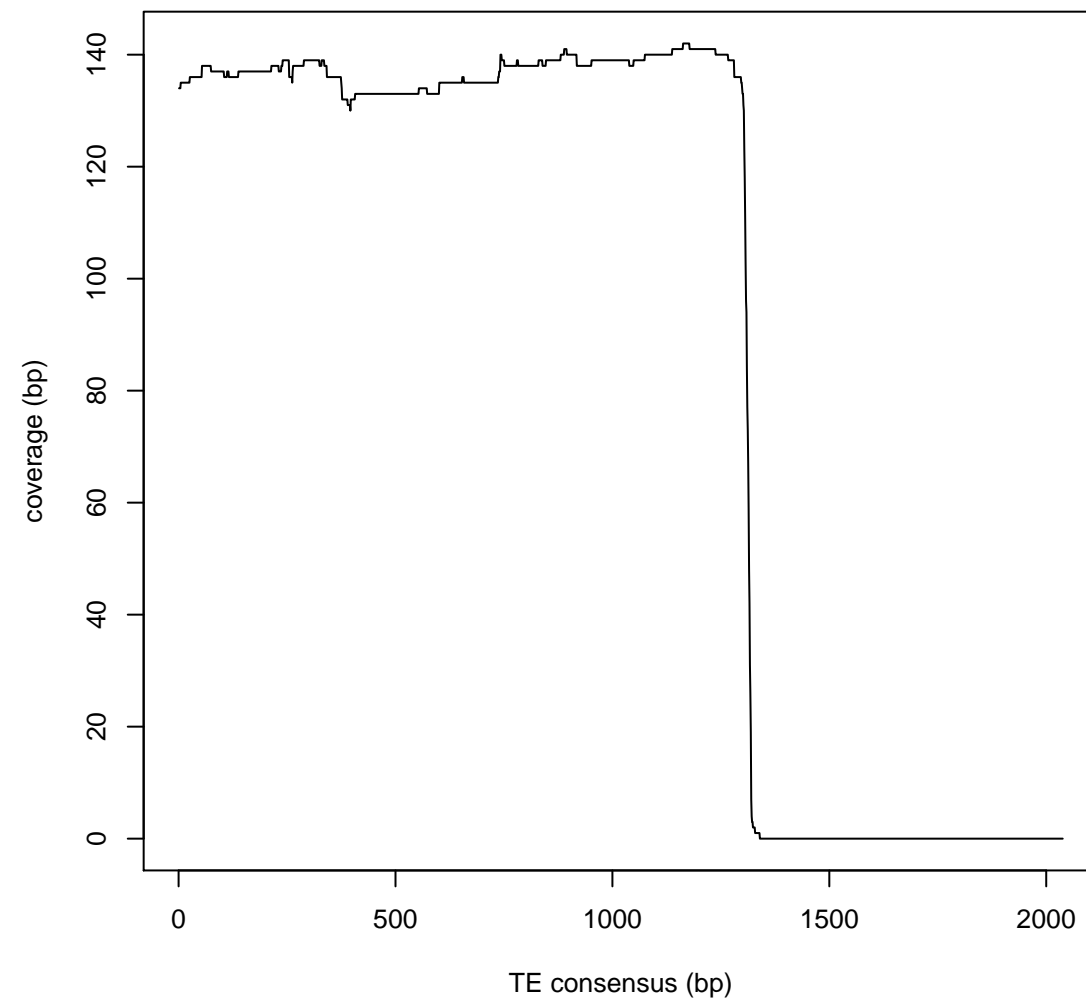


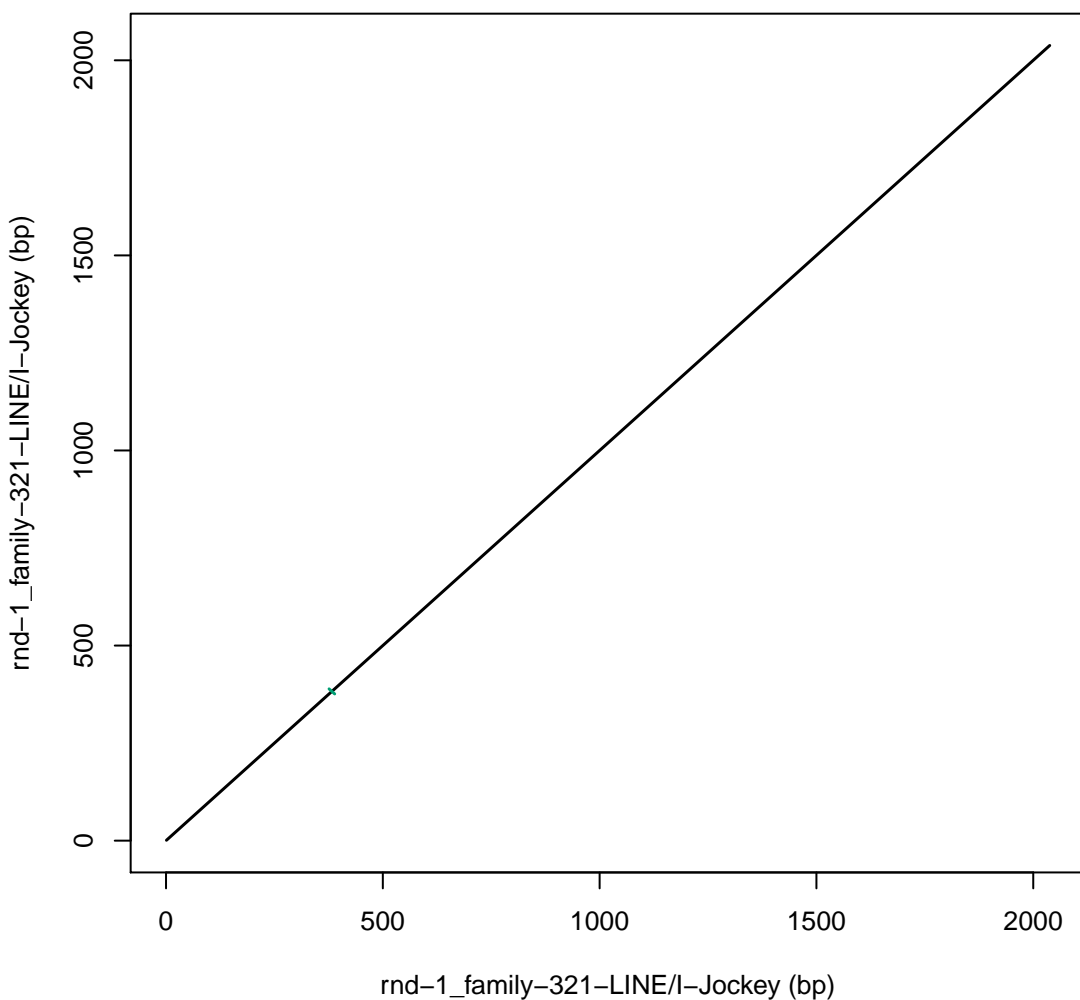
TE: rnd-1_family-321-LINE/I-Jockey
consensus size: 2038bp; fragments: 170; full length: 0 (≥ 1834.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

