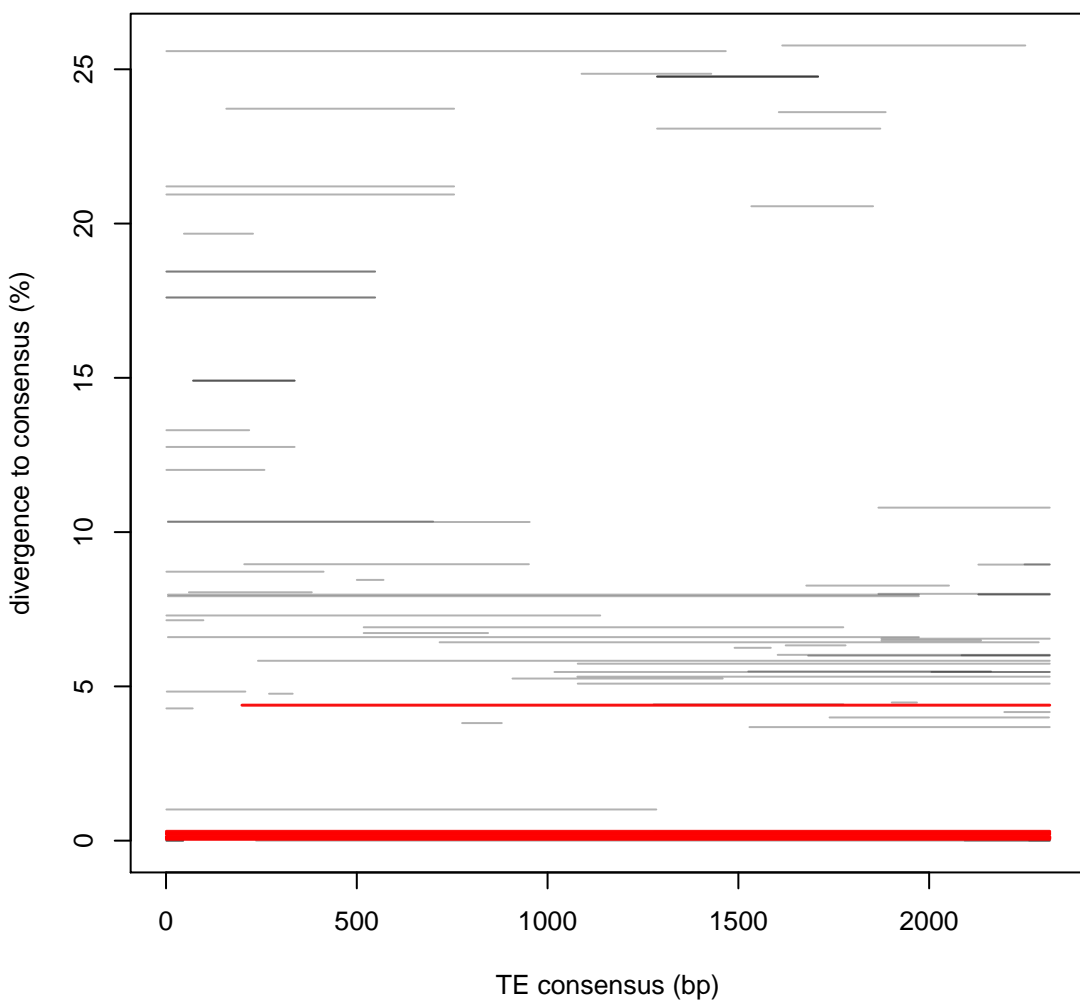
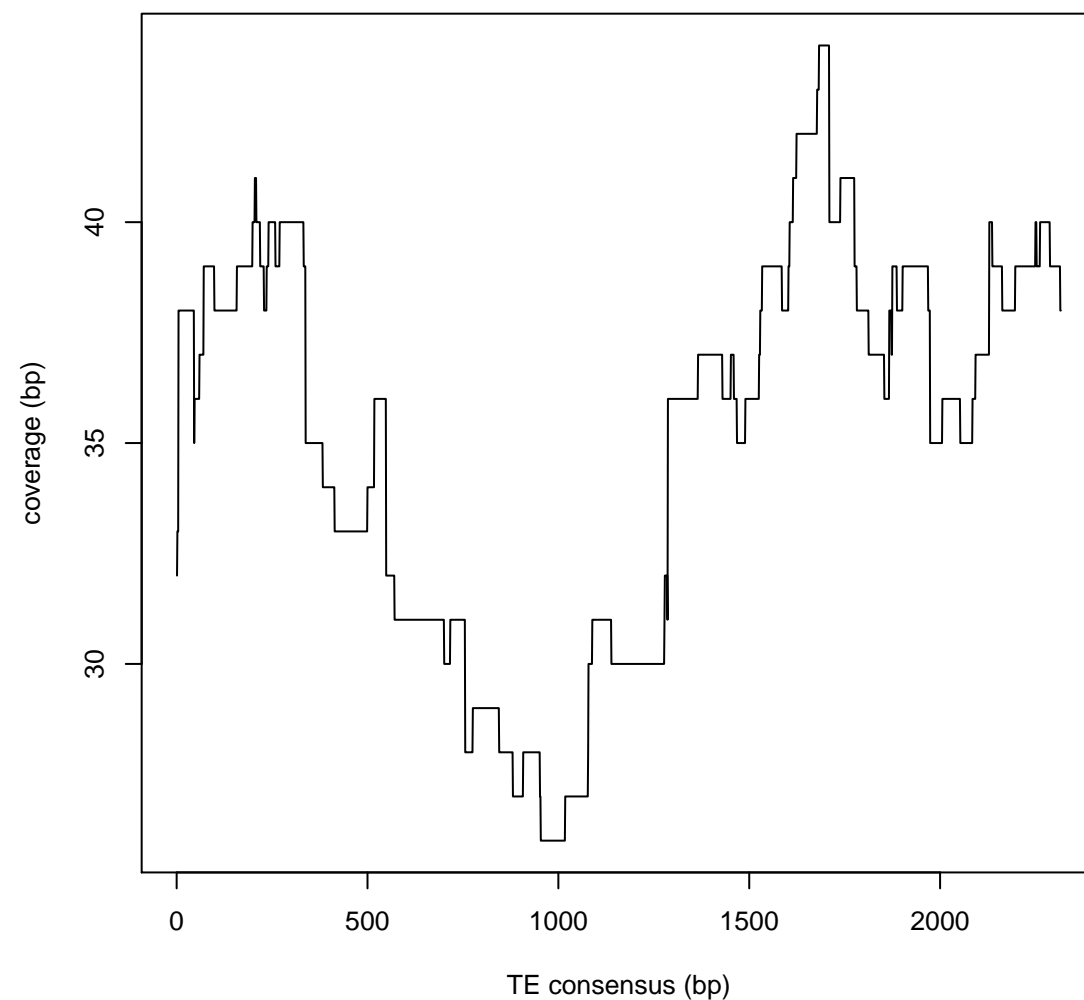


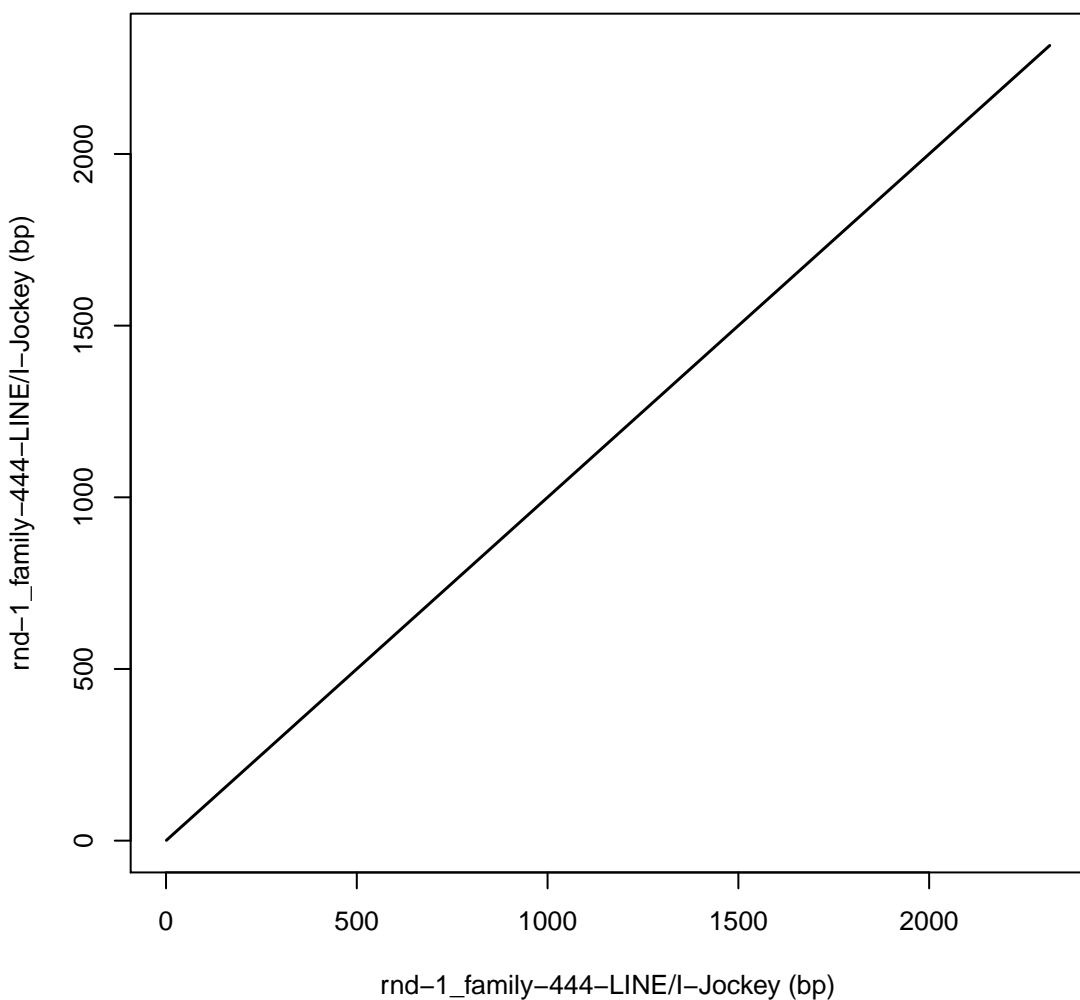
TE: rnd-1_family-444-LINE/I-Jockey
consensus size: 2316bp; fragments: 94; full length: 14 (≥ 2084.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

