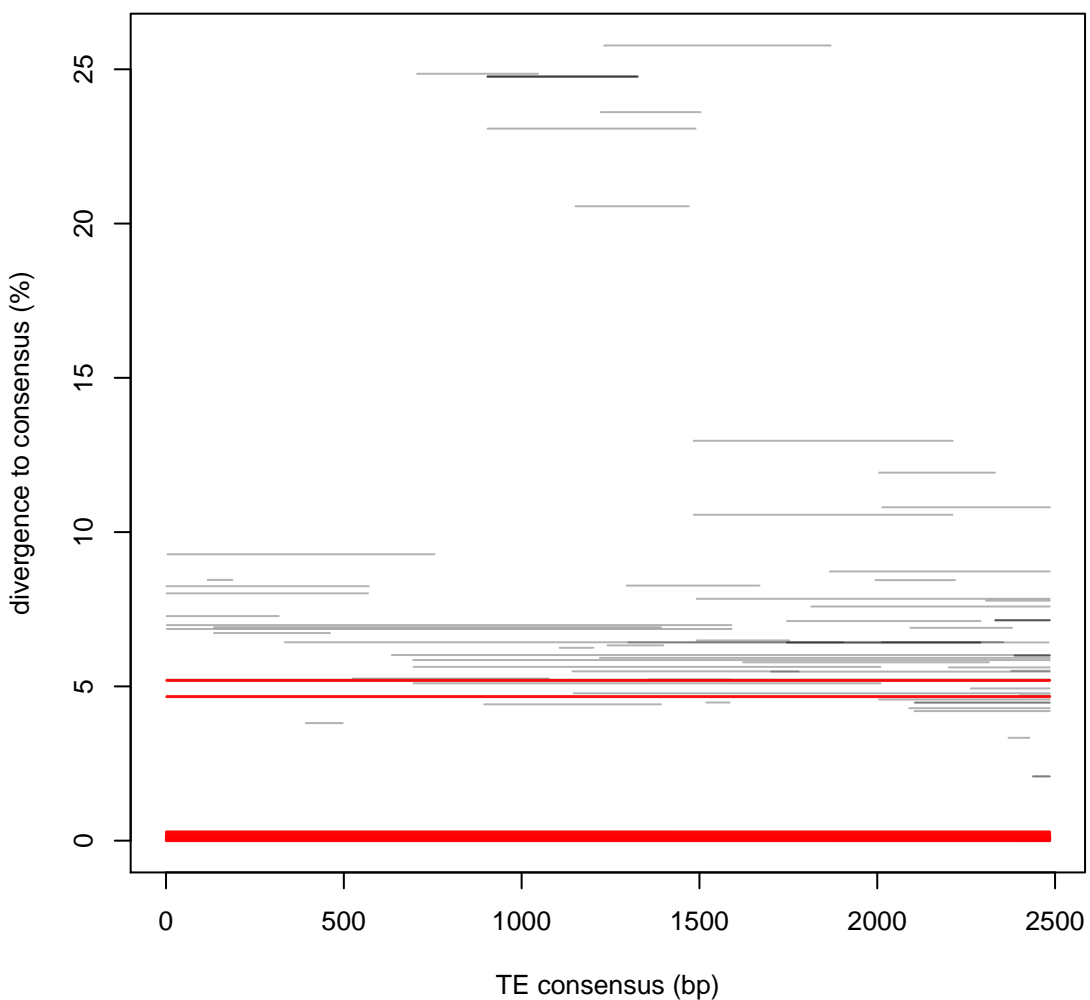
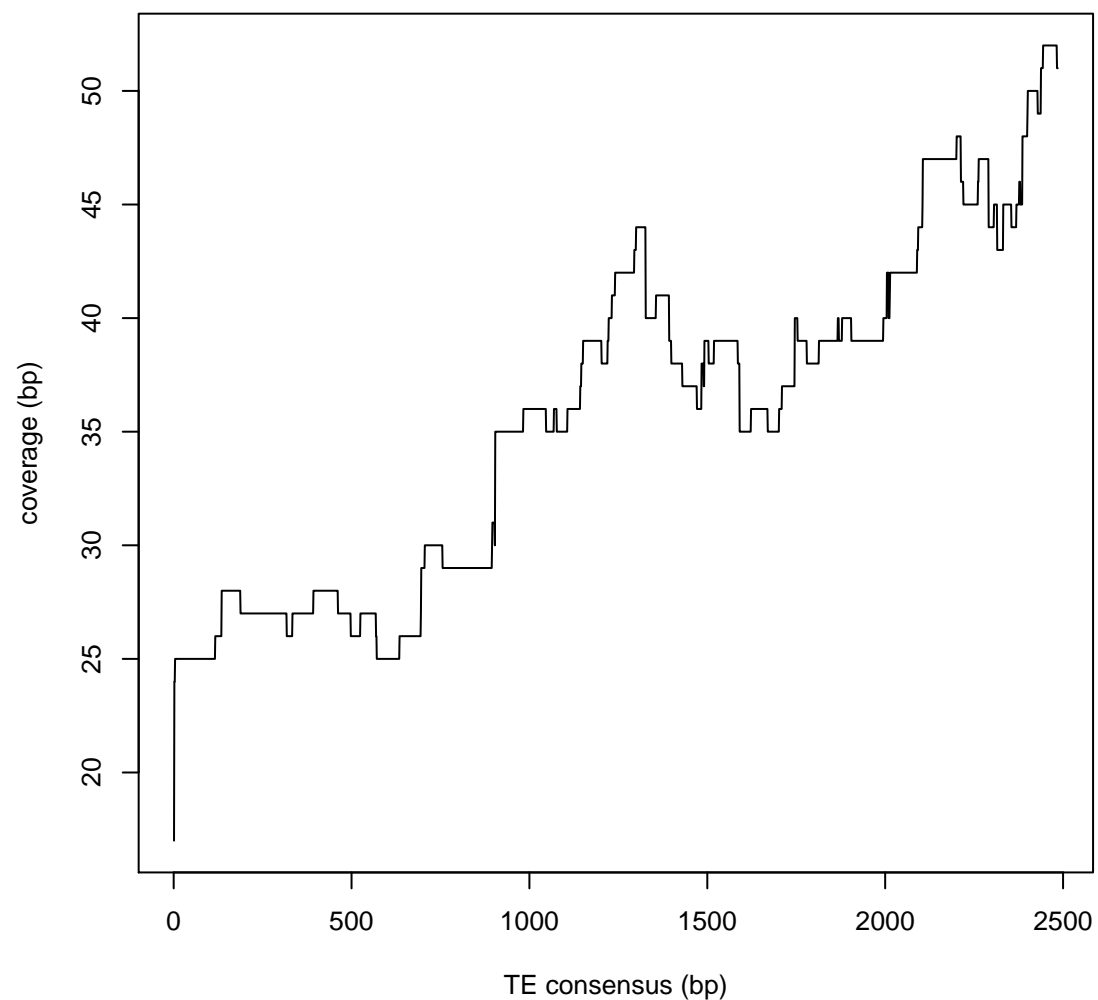


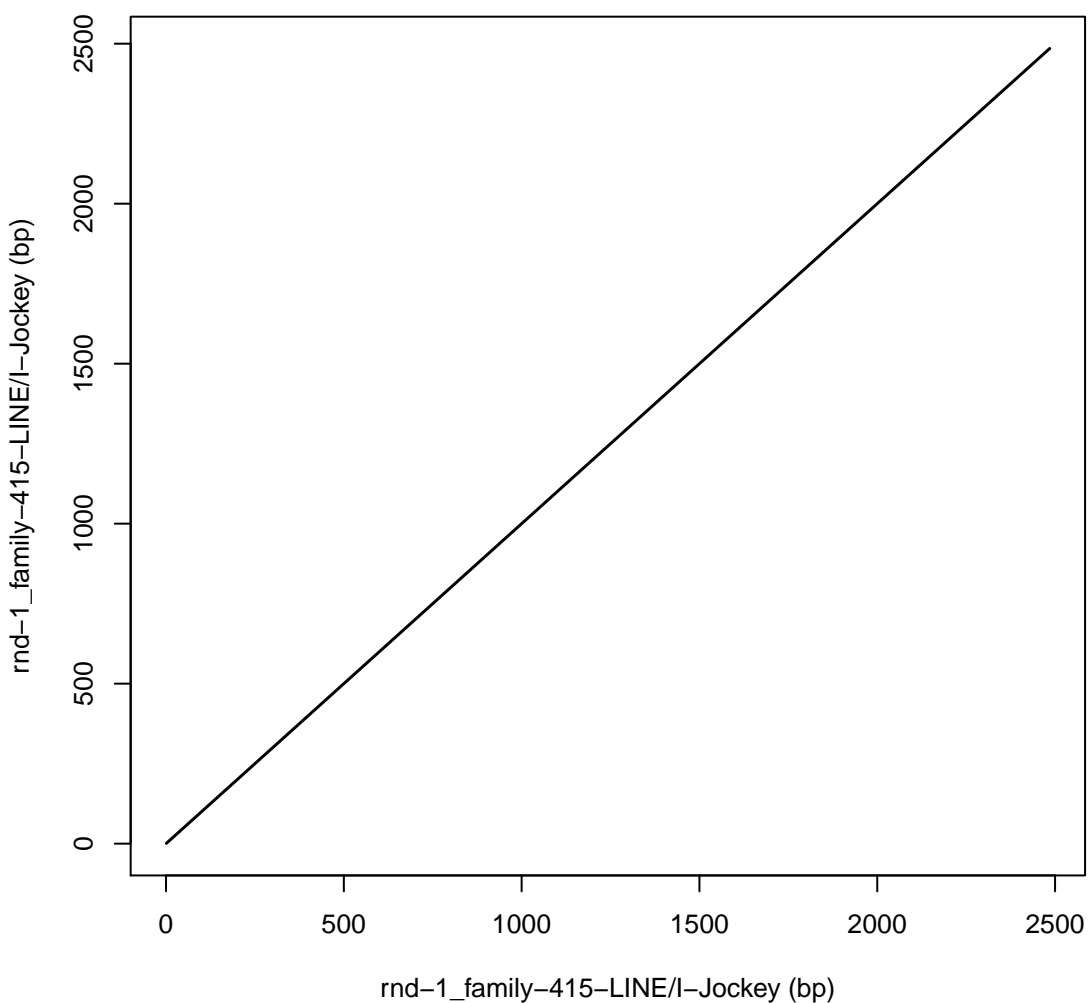
TE: rnd-1_family-415-LINE/I-Jockey
consensus size: 2485bp; fragments: 97; full length: 16 (≥ 2236.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

