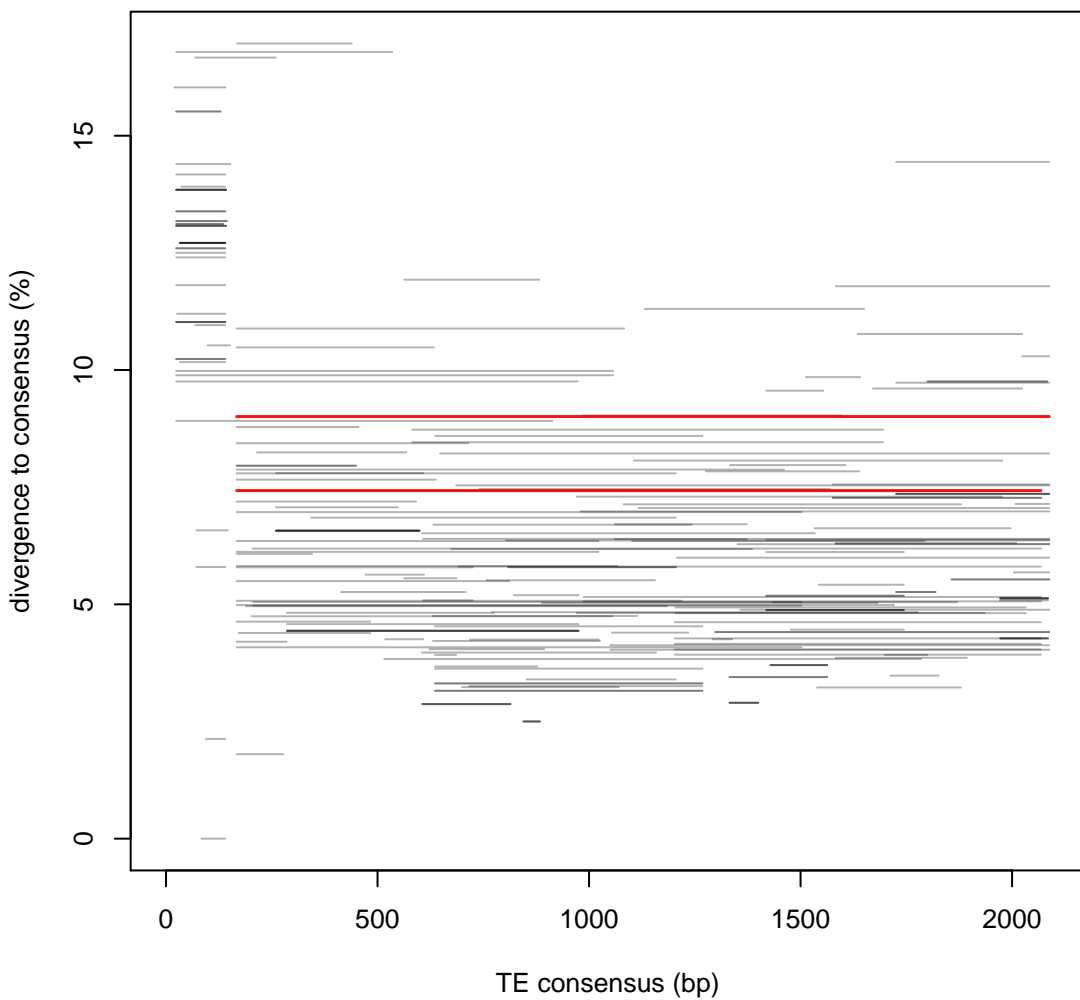
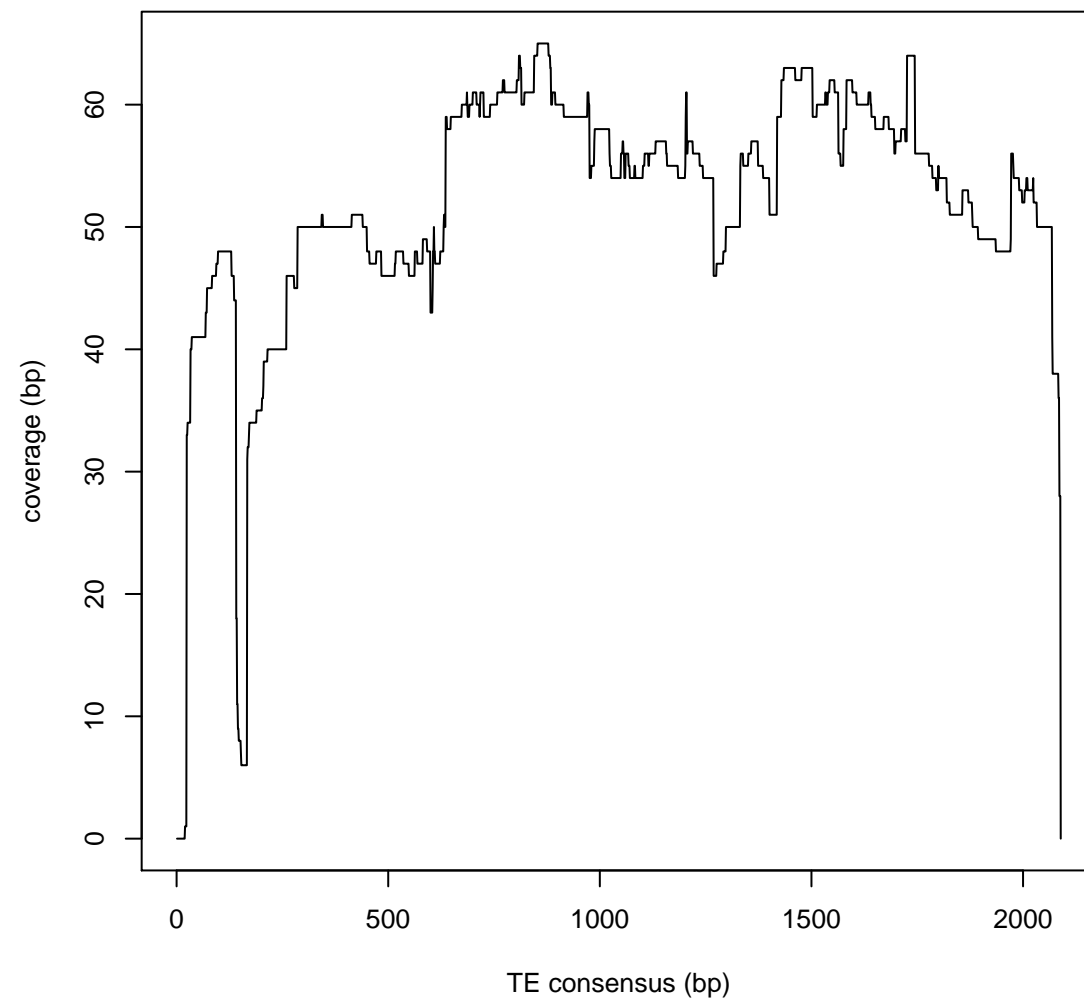


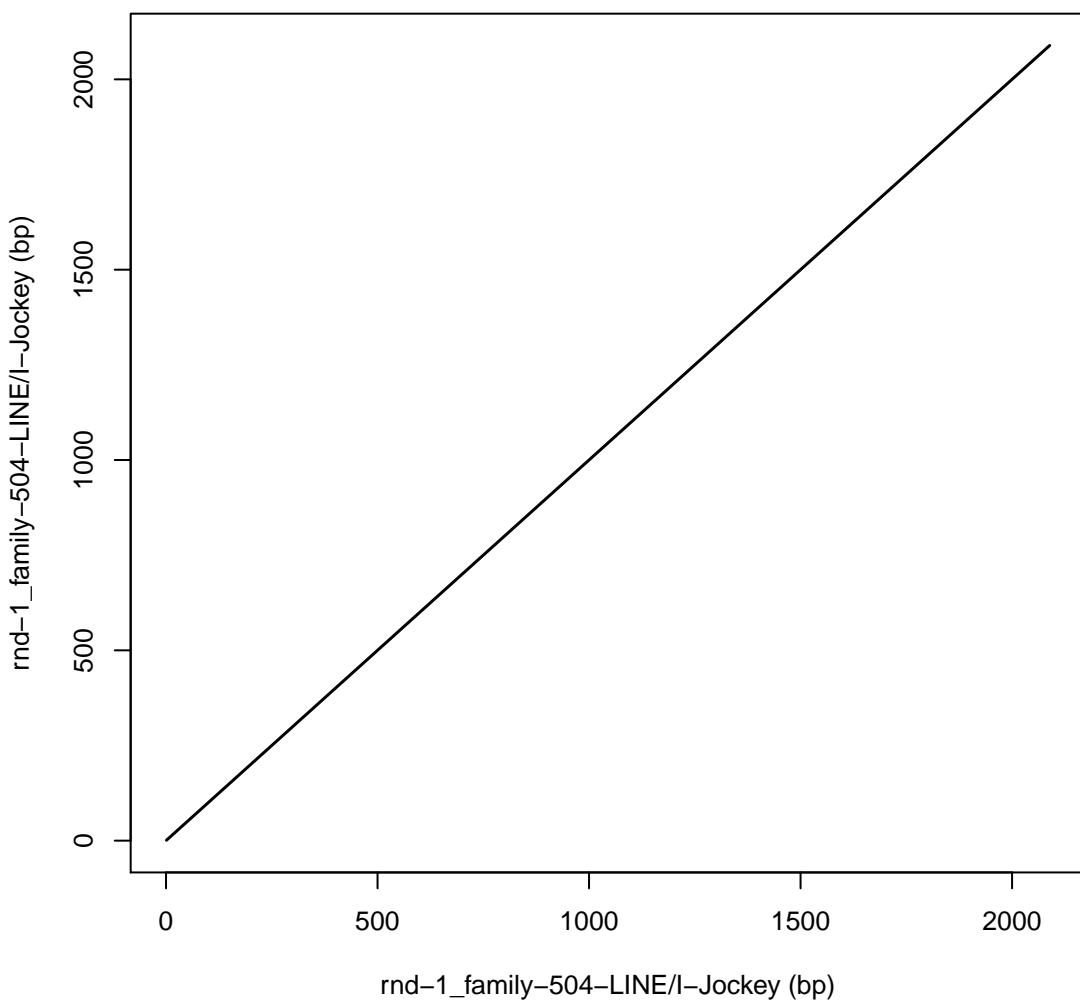
TE: rnd-1_family-504-LINE/I-Jockey
consensus size: 2089bp; fragments: 229; full length: 2 (≥ 1880.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

