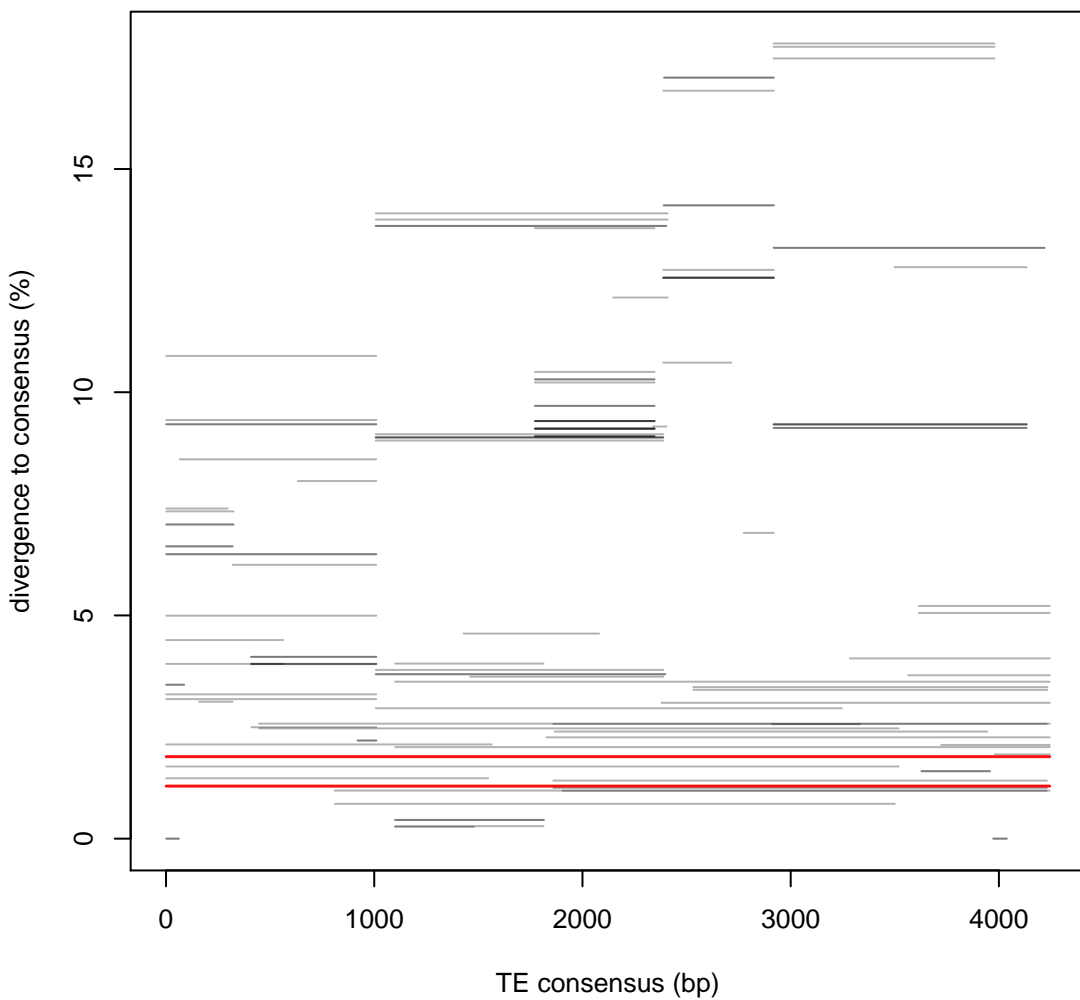
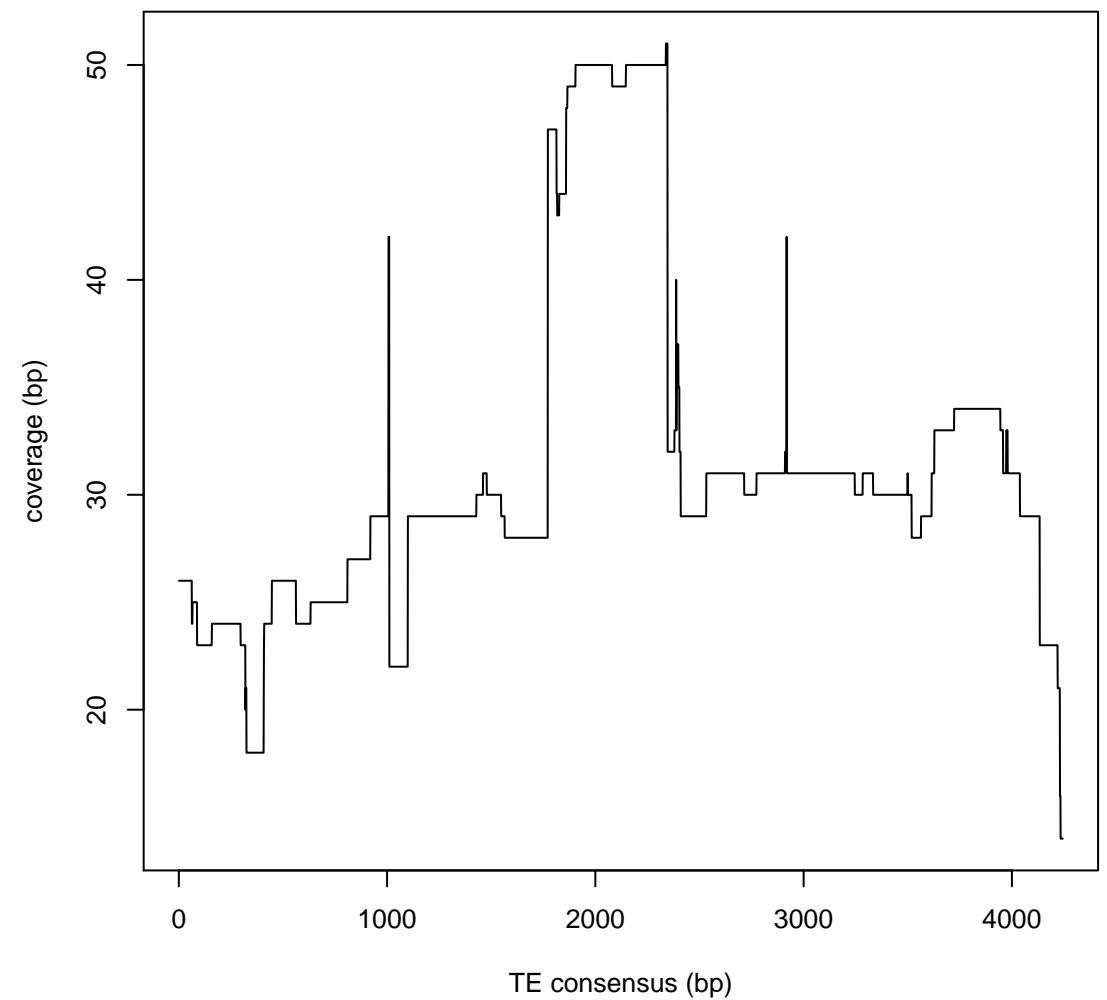


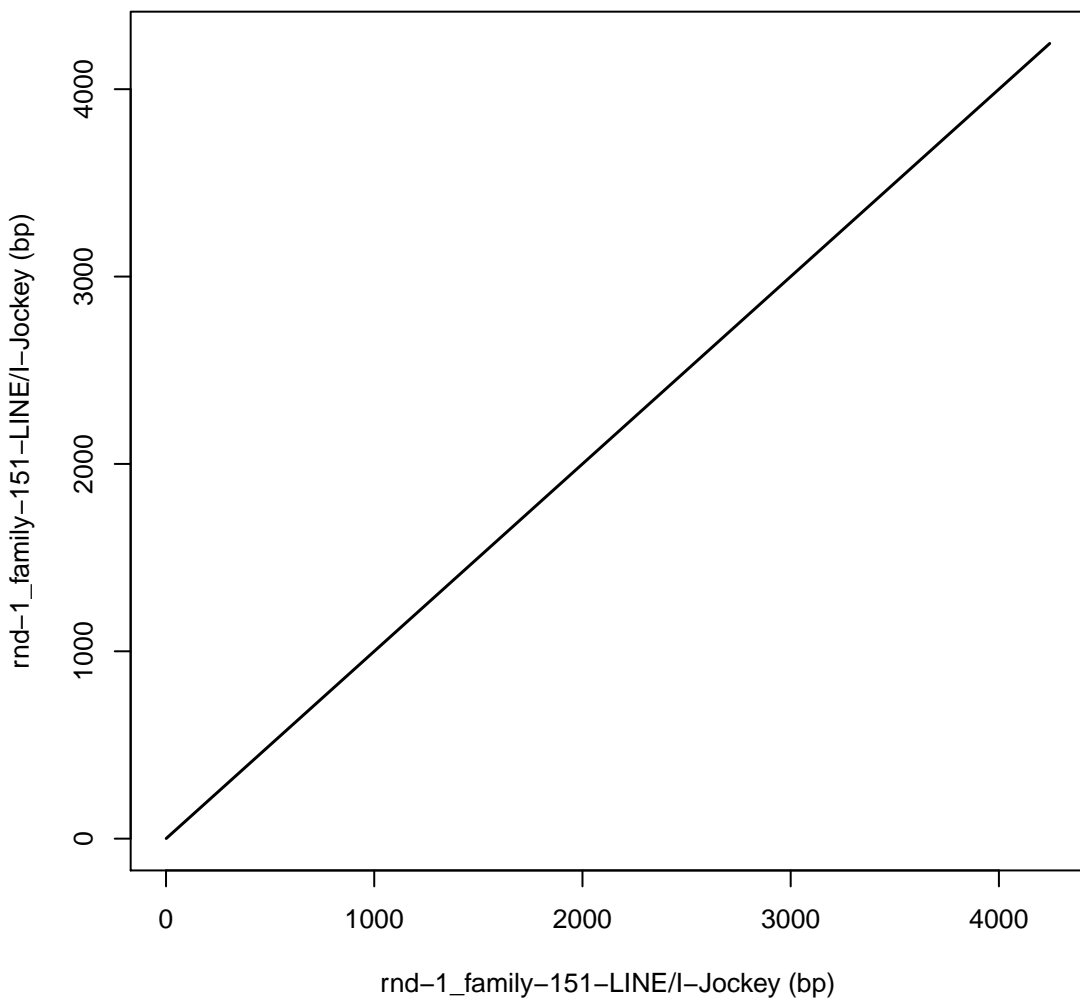
TE: rnd-1_family-151-LINE/I-Jockey
consensus size: 4244bp; fragments: 129; full length: 2 (≥ 3819.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

