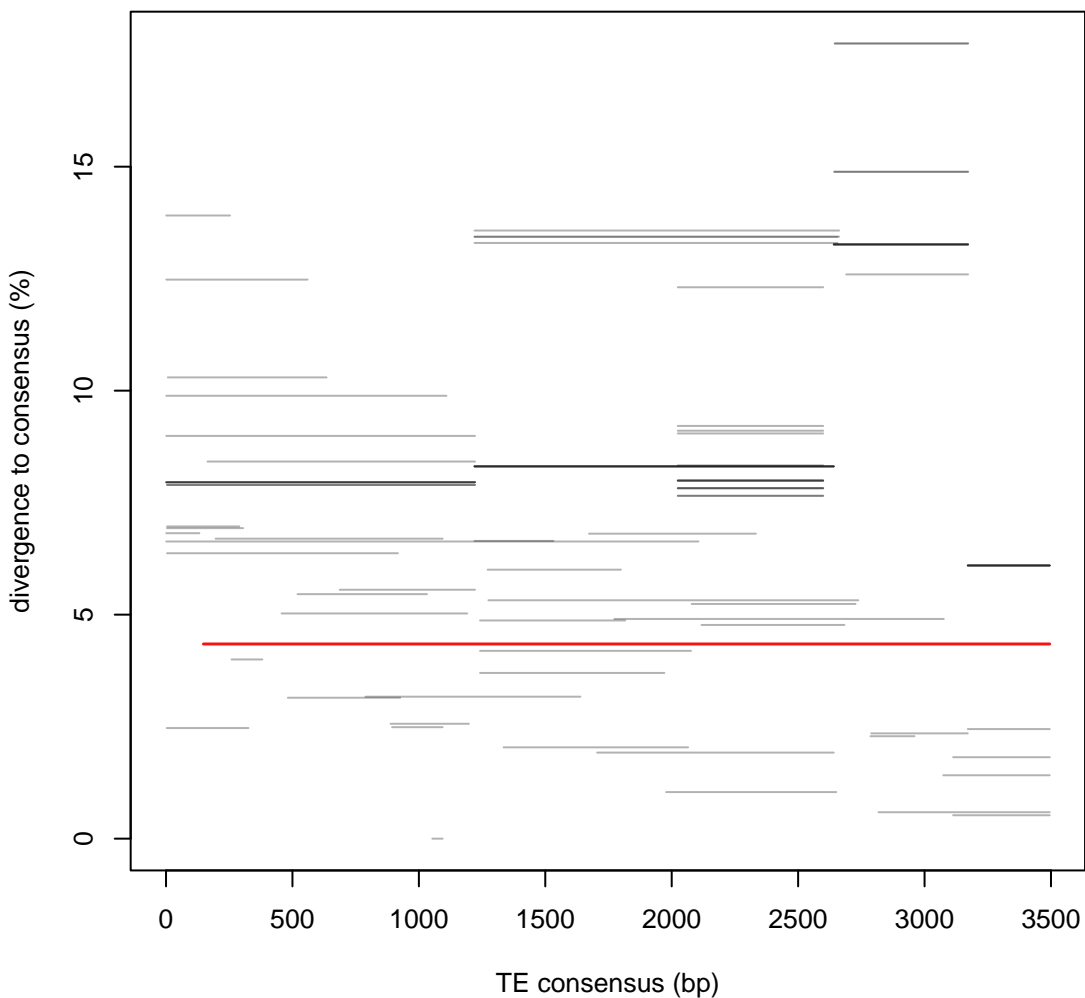
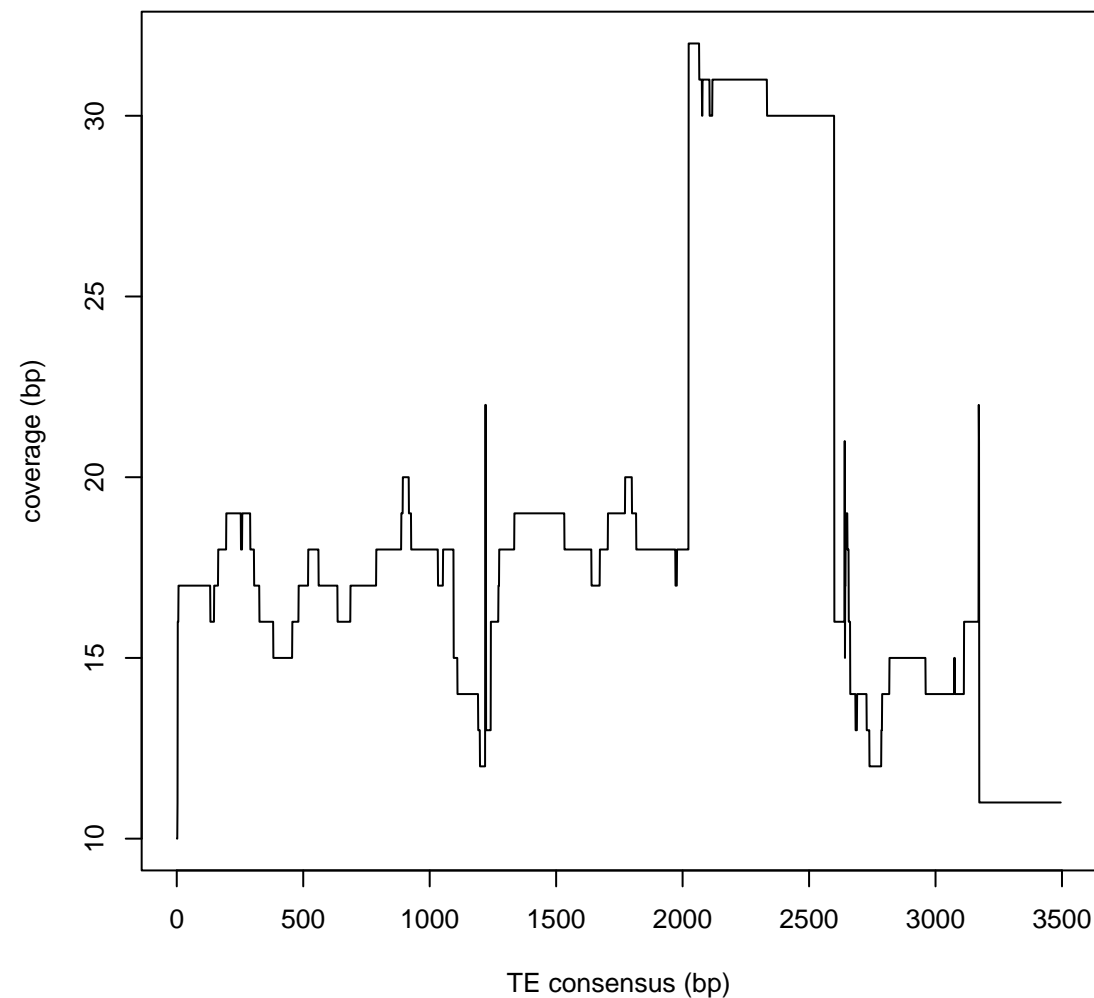


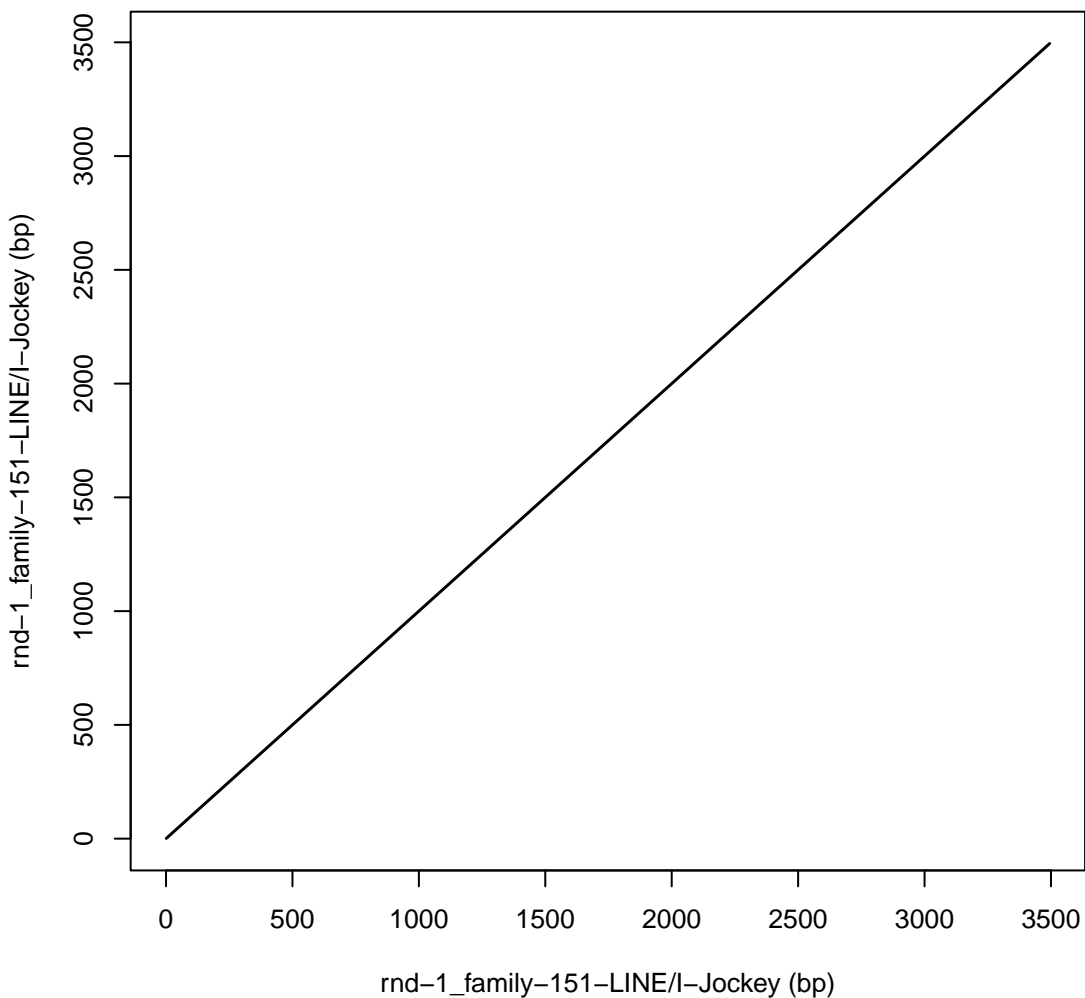
TE: rnd-1_family-151-LINE/I-Jockey
consensus size: 3495bp; fragments: 87; full length: 1 (>=3145.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

no orf >400 bp detected