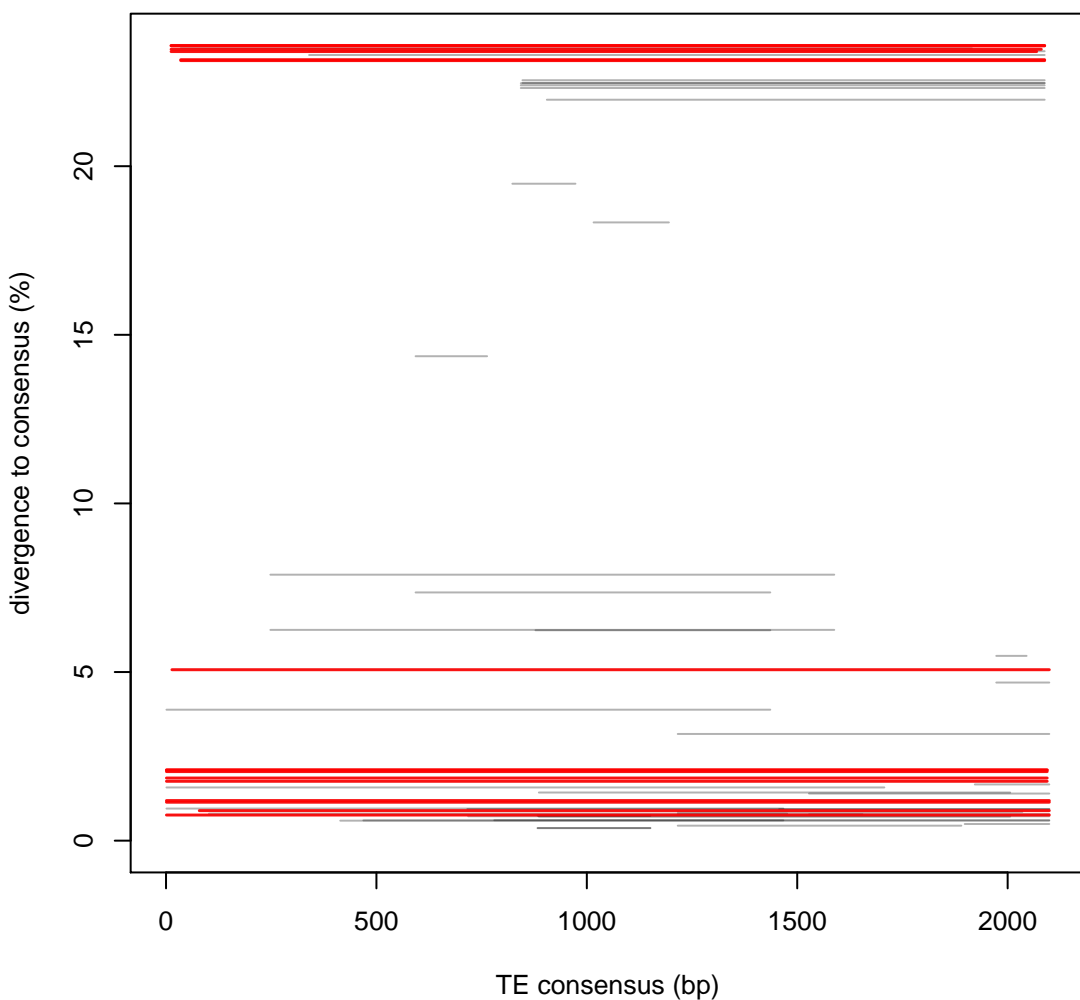
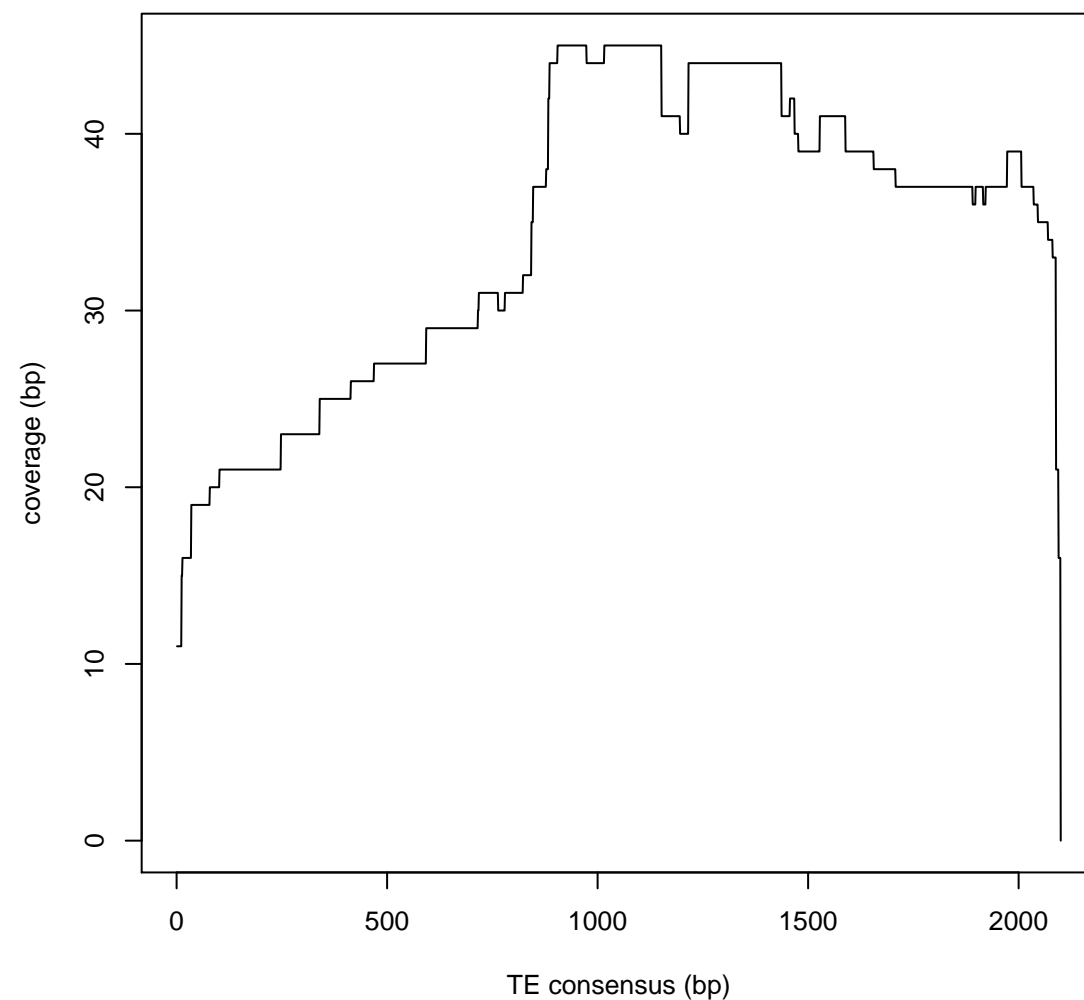


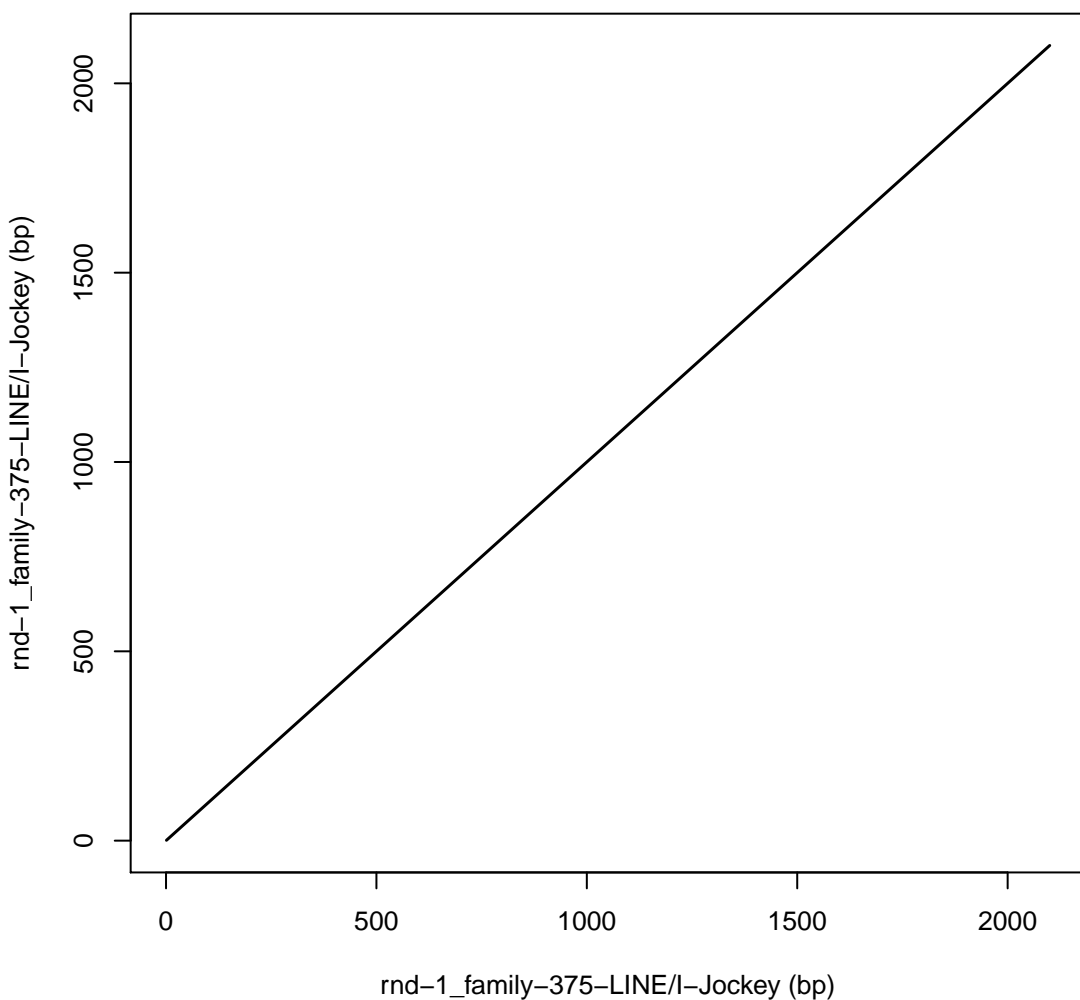
TE: rnd-1_family-375-LINE/I-Jockey
consensus size: 2100bp; fragments: 58; full length: 16 (≥ 1890 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

