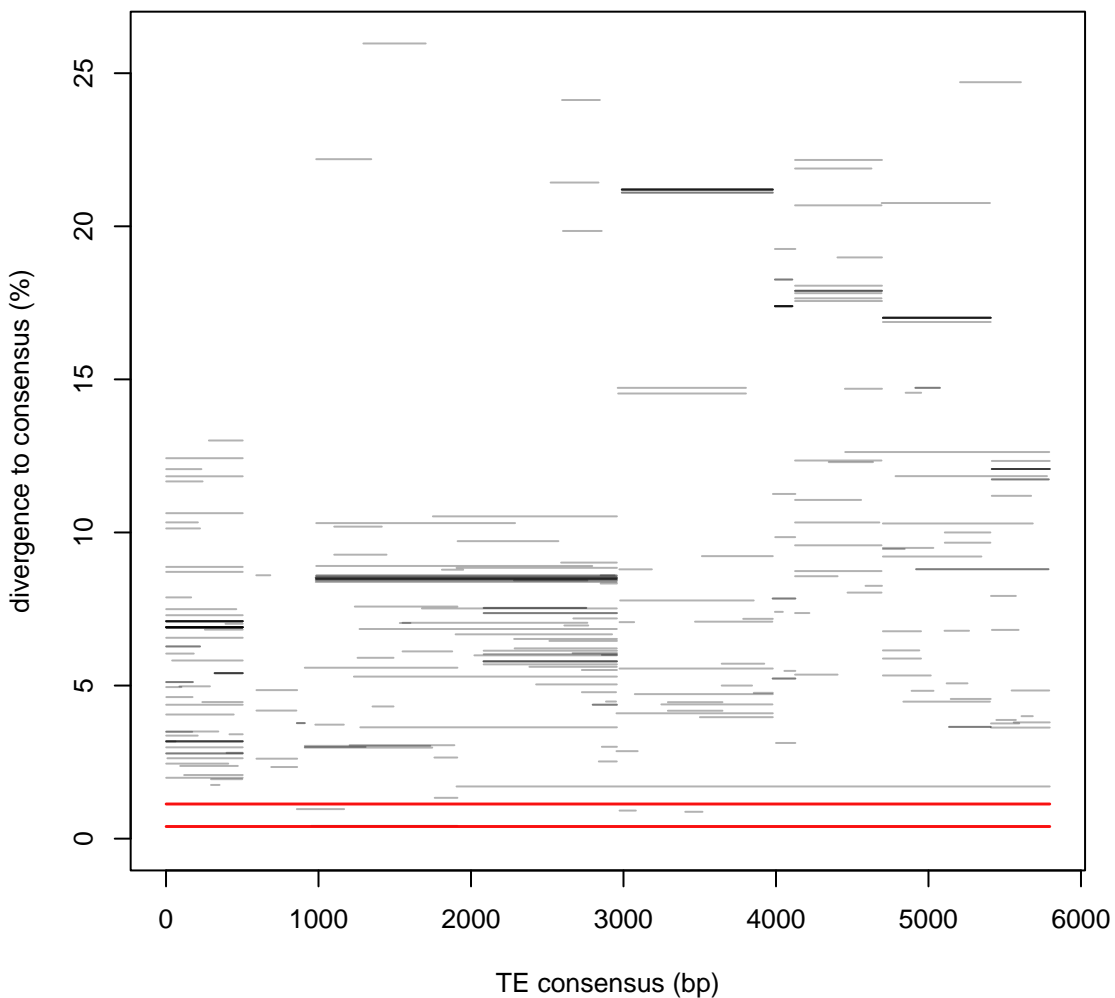
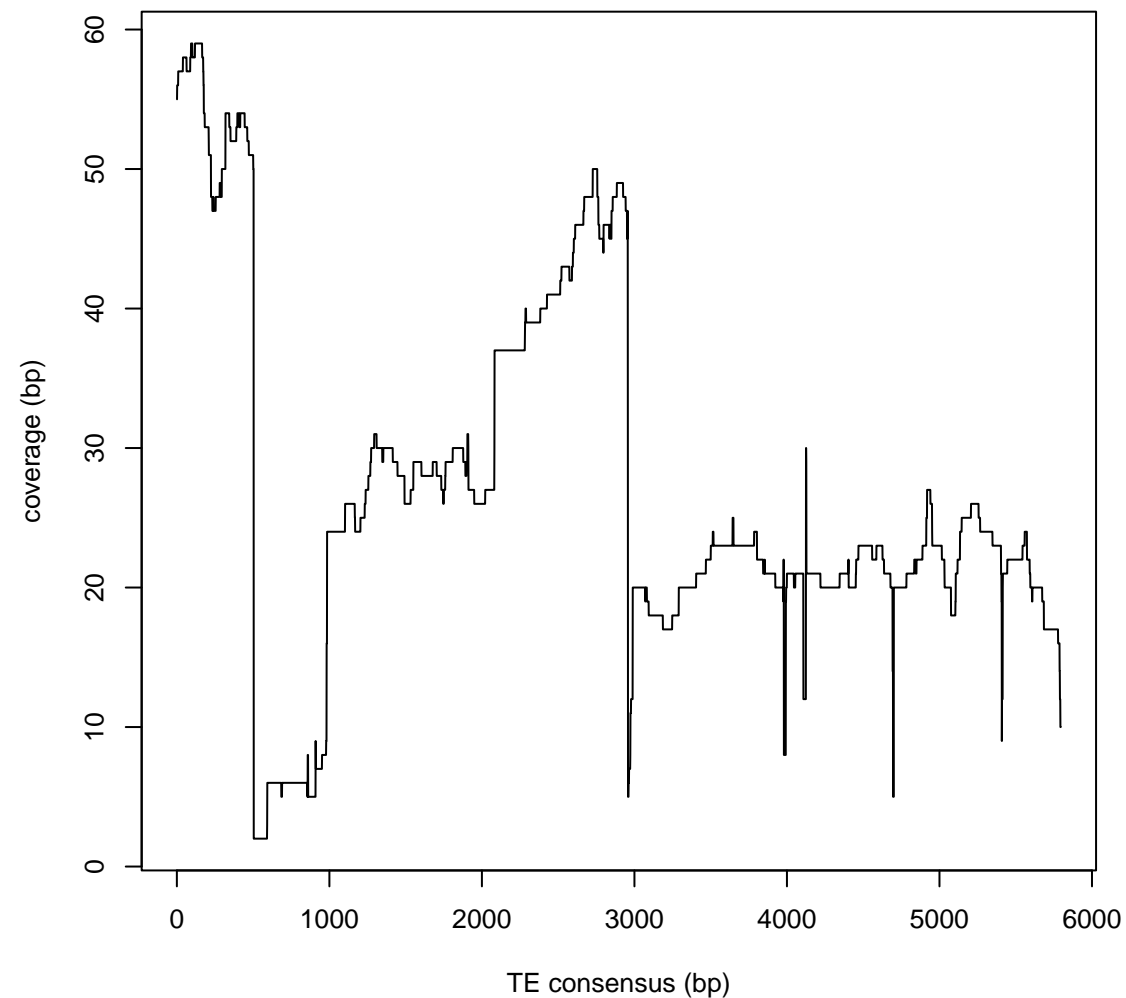


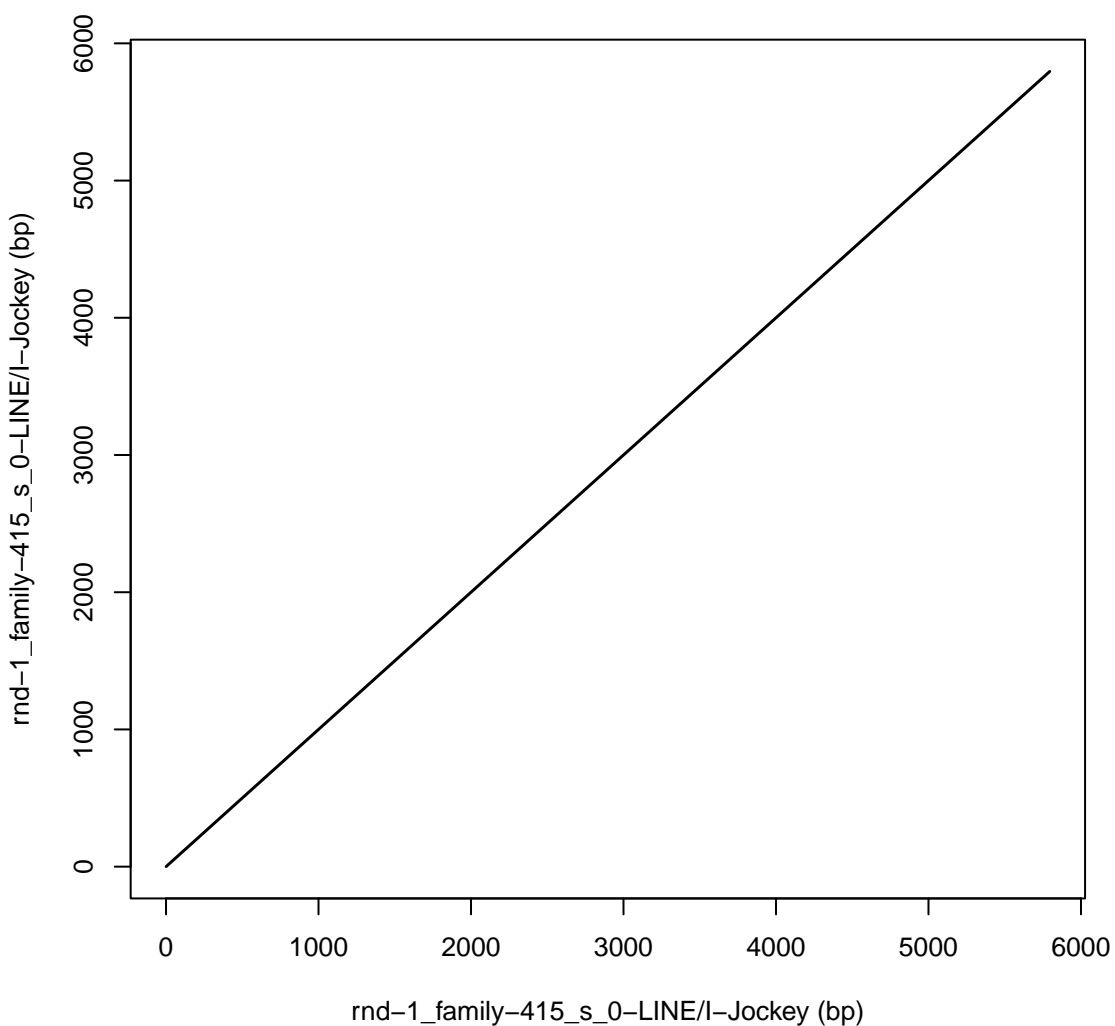
TE: rnd-1_family-415_s_0-LINE/I-Jockey
consensus size: 5795bp; fragments: 278; full length: 2 (>=5215.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

