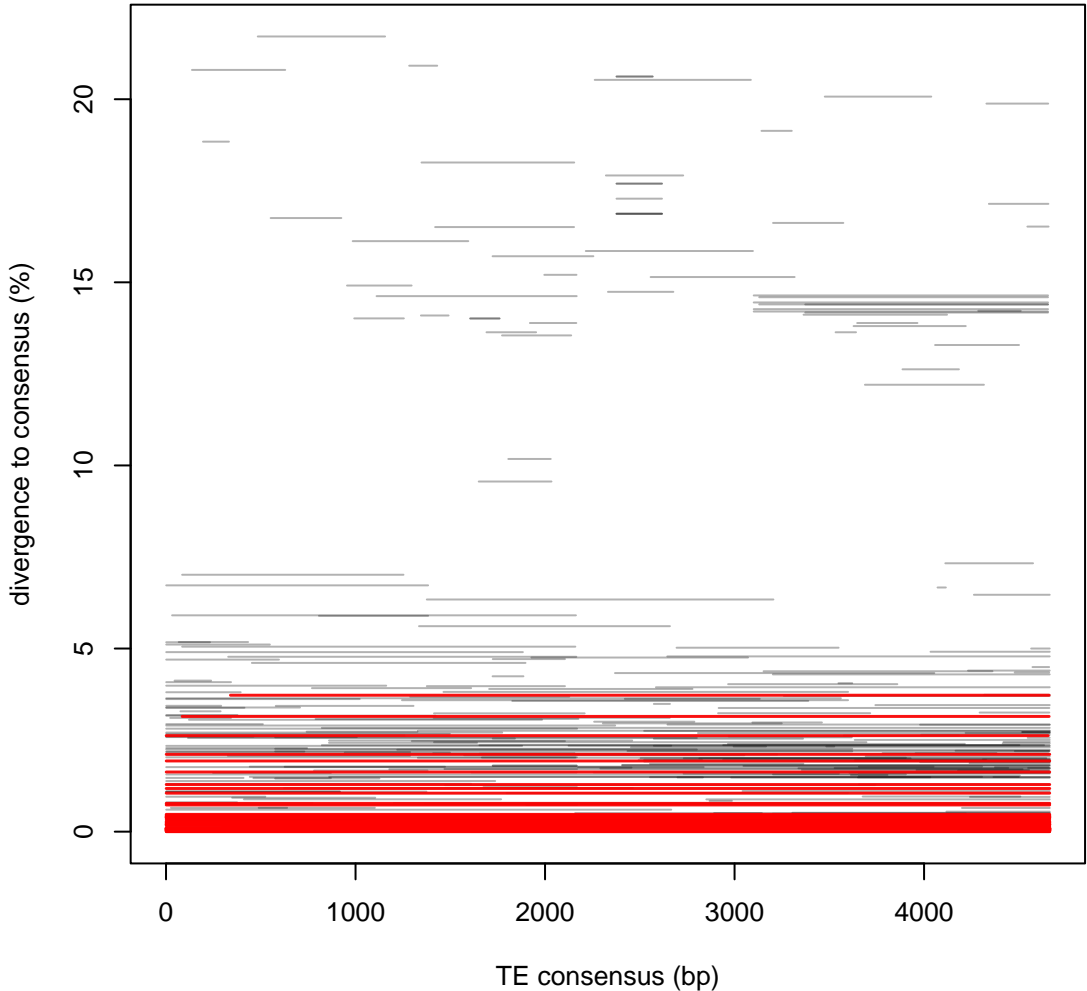
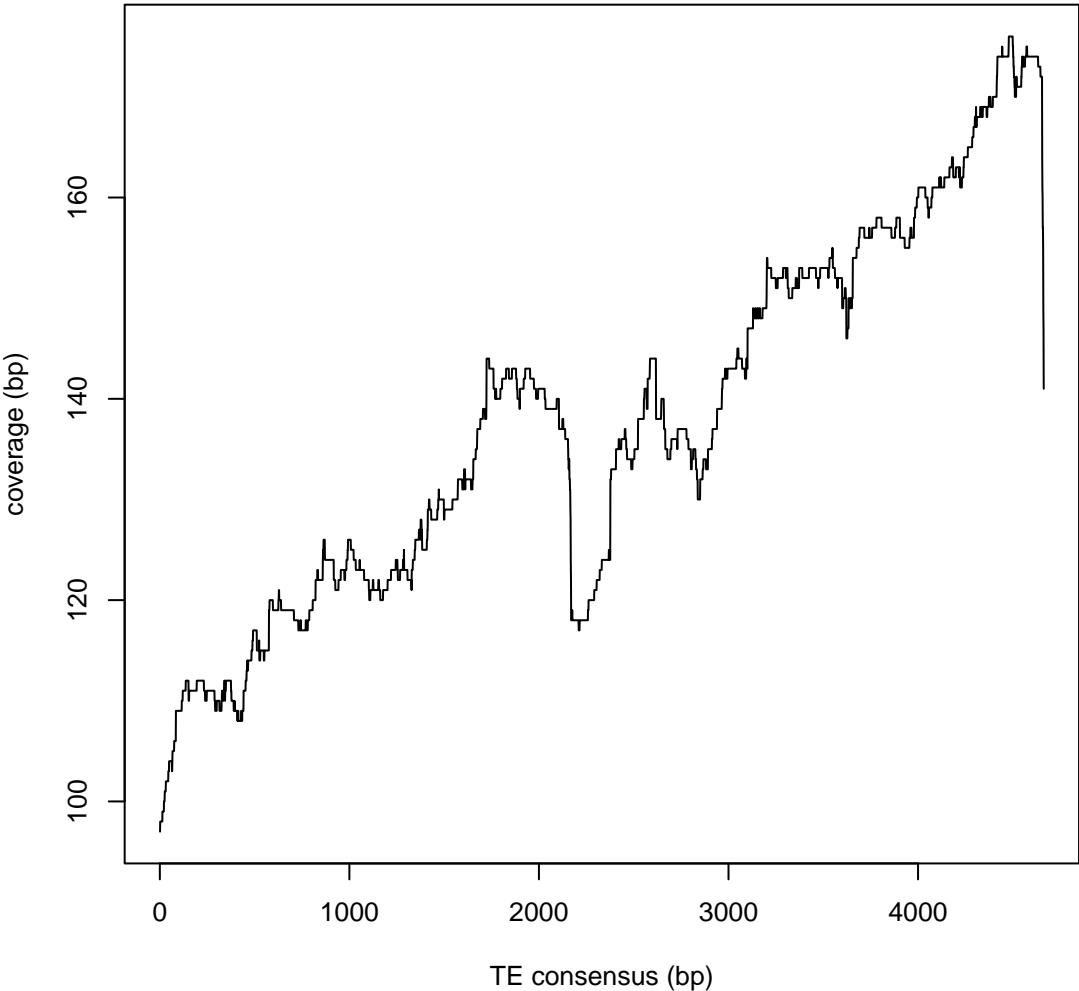


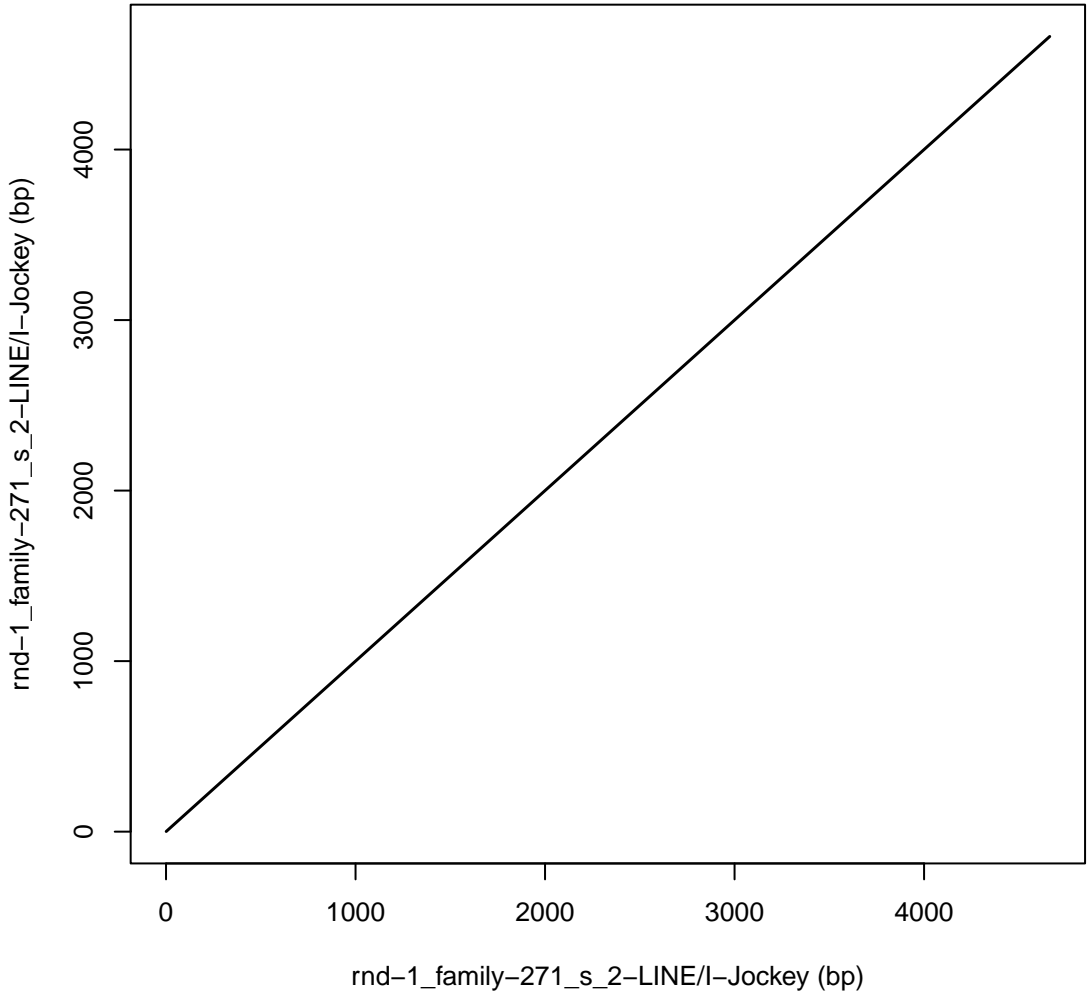
TE: rnd-1_family-271_s_2-LINE/I-Jockey
consensus size: 4663bp; fragments: 392; full length: 53 (>=4196.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

