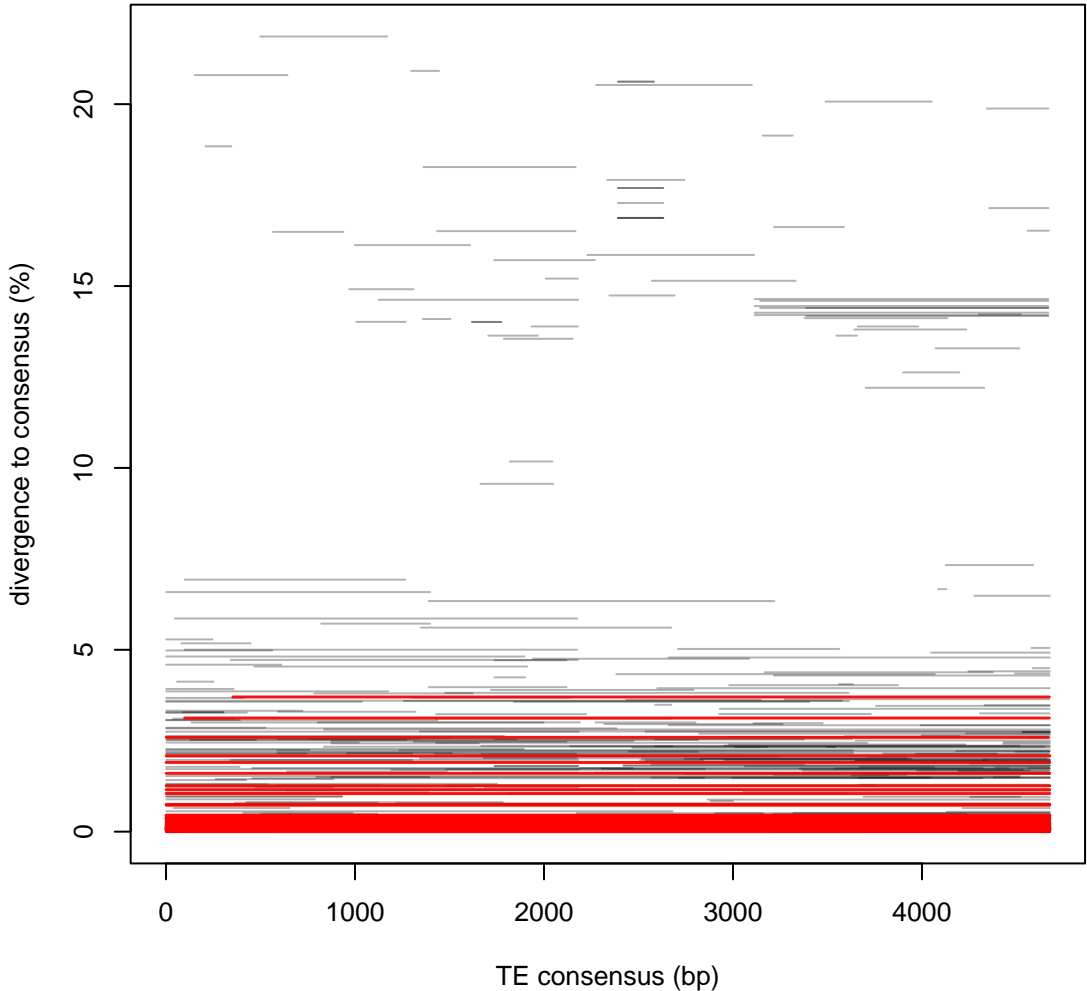
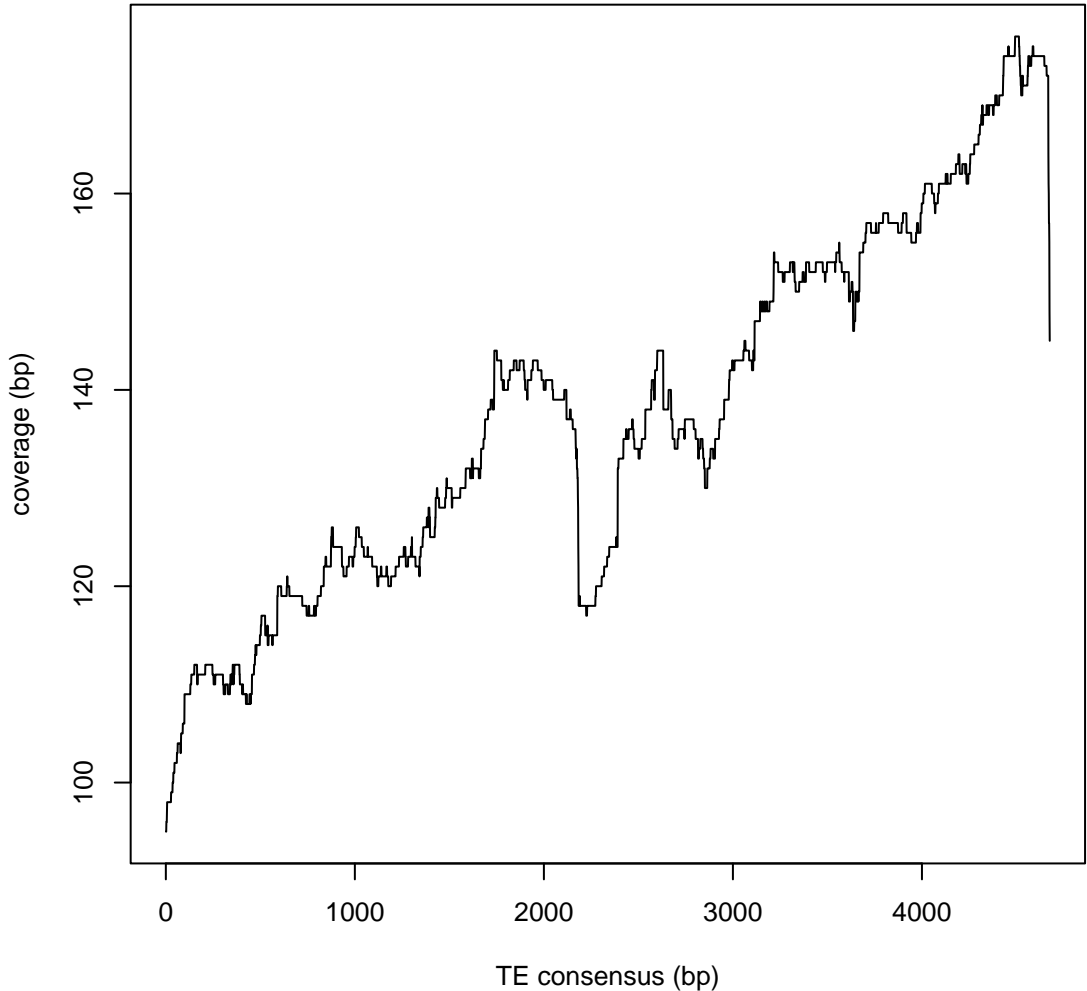


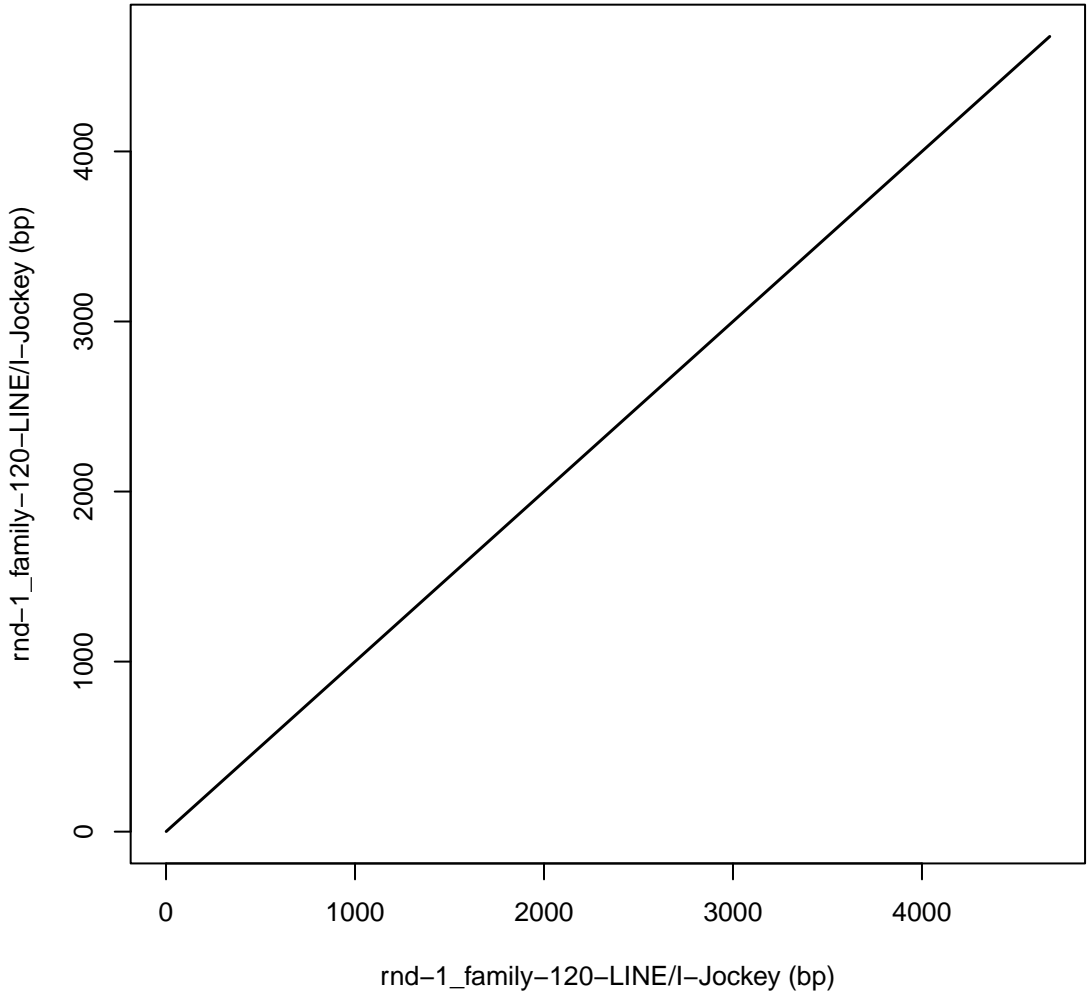
TE: rnd-1_family-120-LINE/I-Jockey
consensus size: 4676bp; fragments: 392; full length: 51 (>=4208.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

