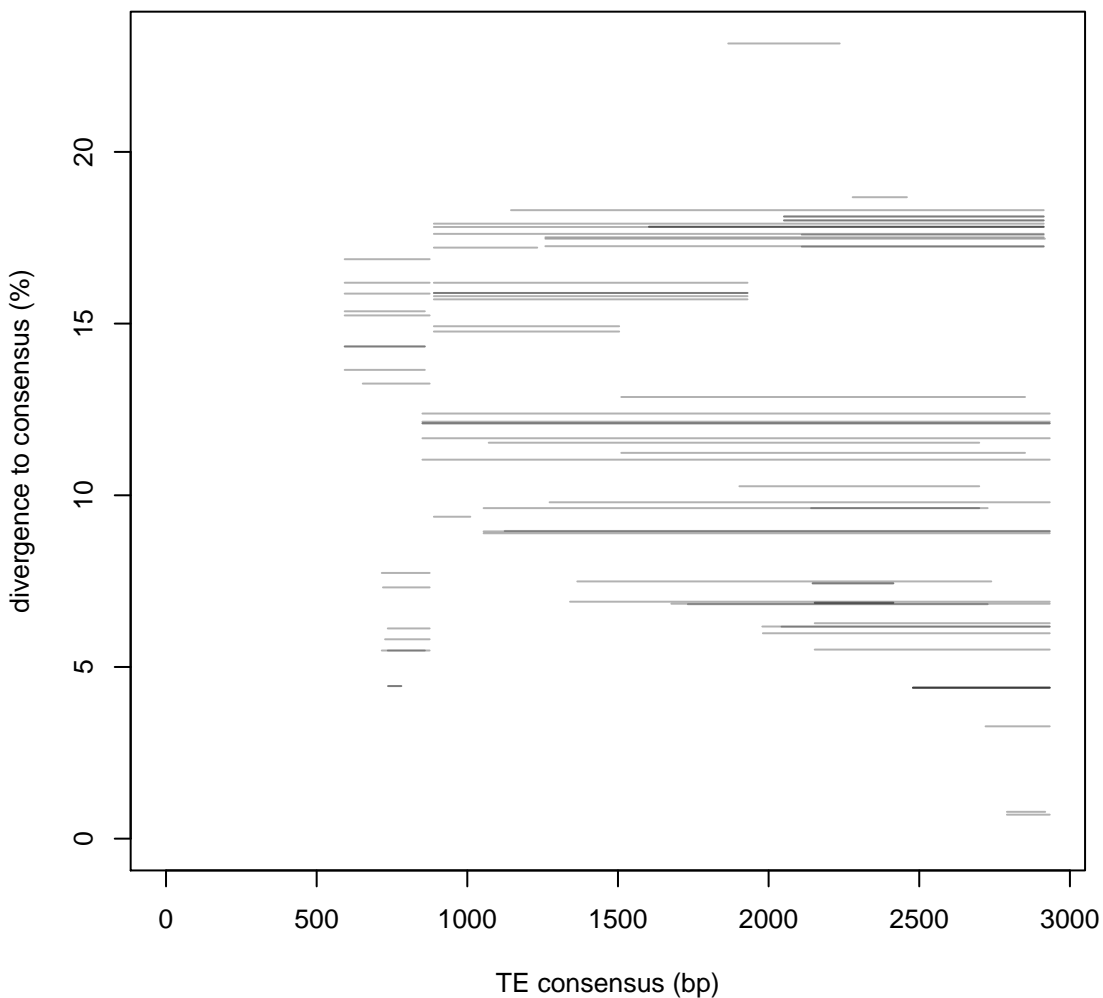
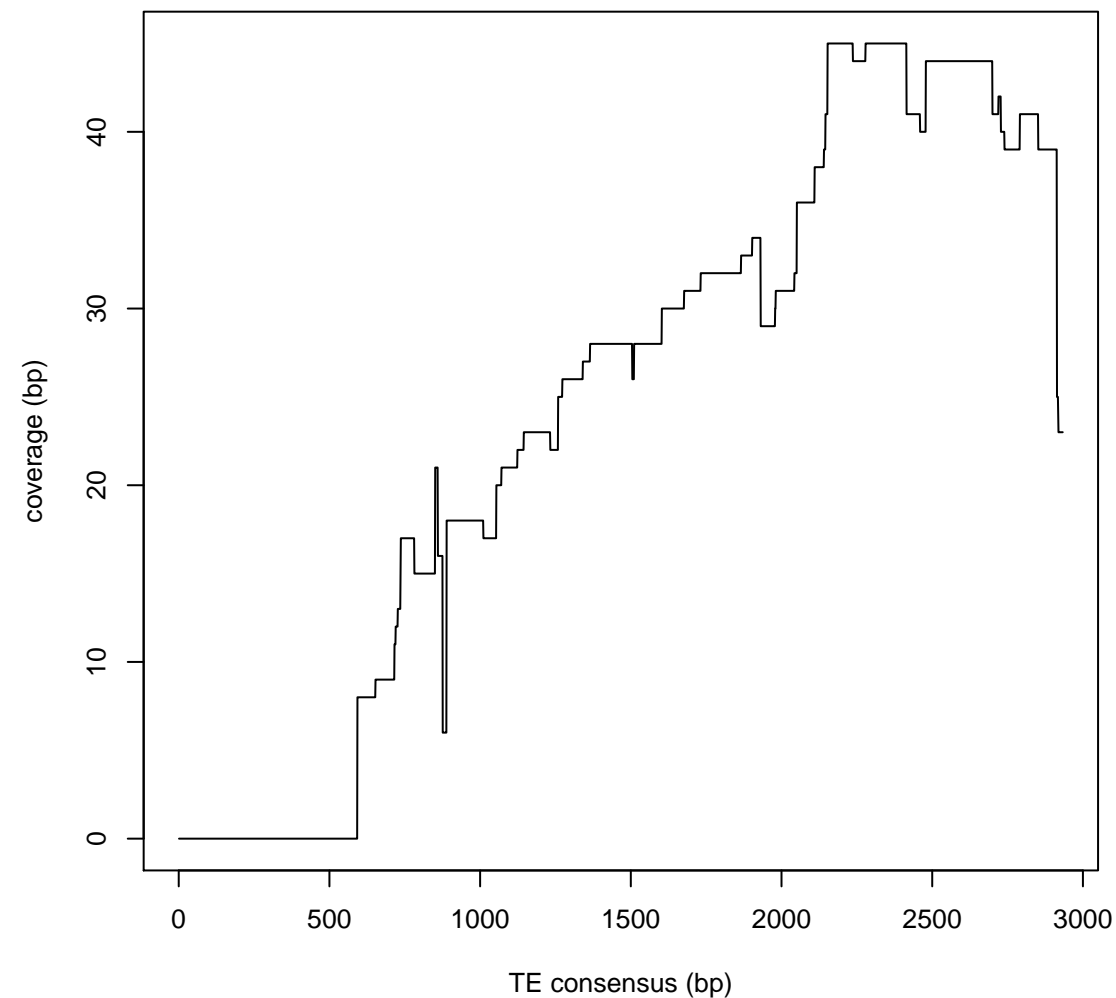


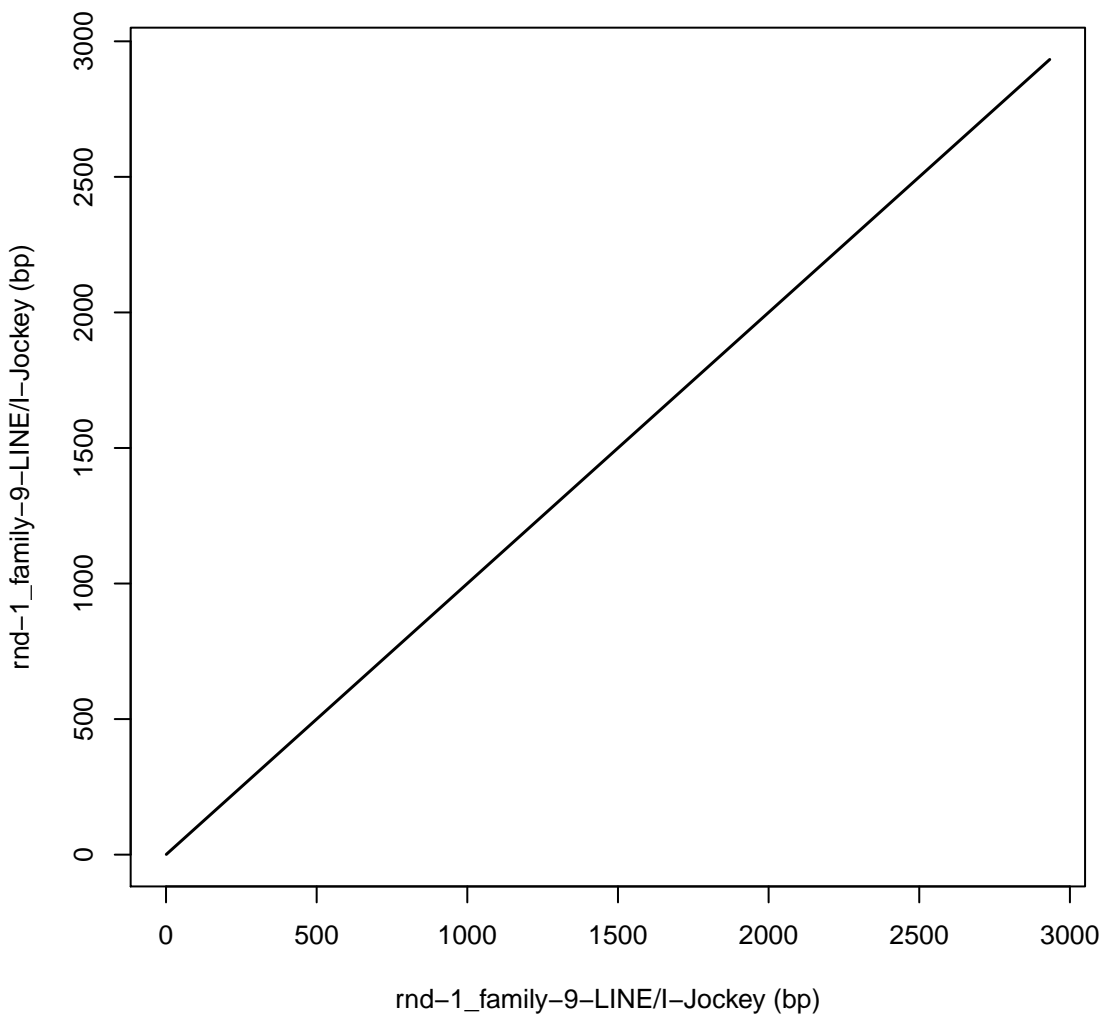
TE: rnd-1_family-9-LINE/I-Jockey
consensus size: 2933bp; fragments: 79; full length: 0 (>=2639.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

