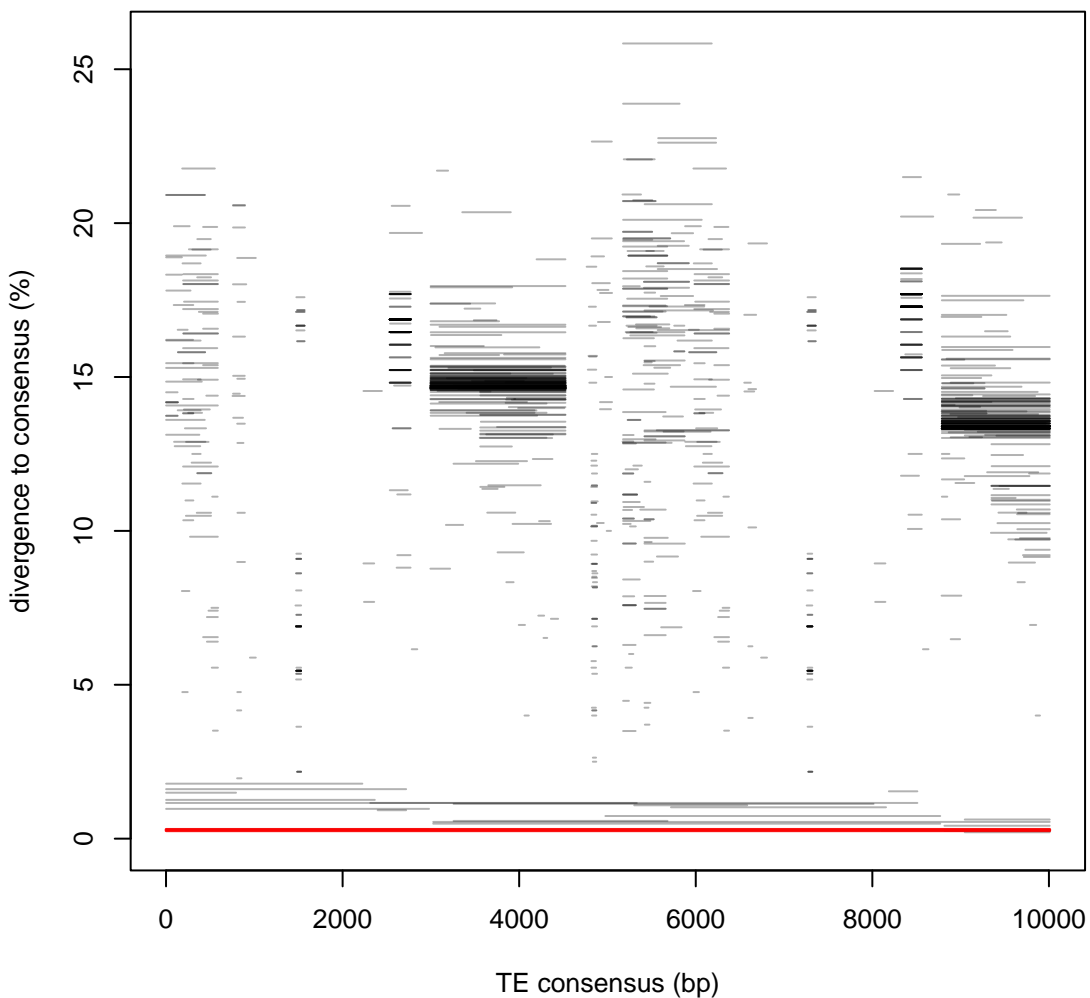
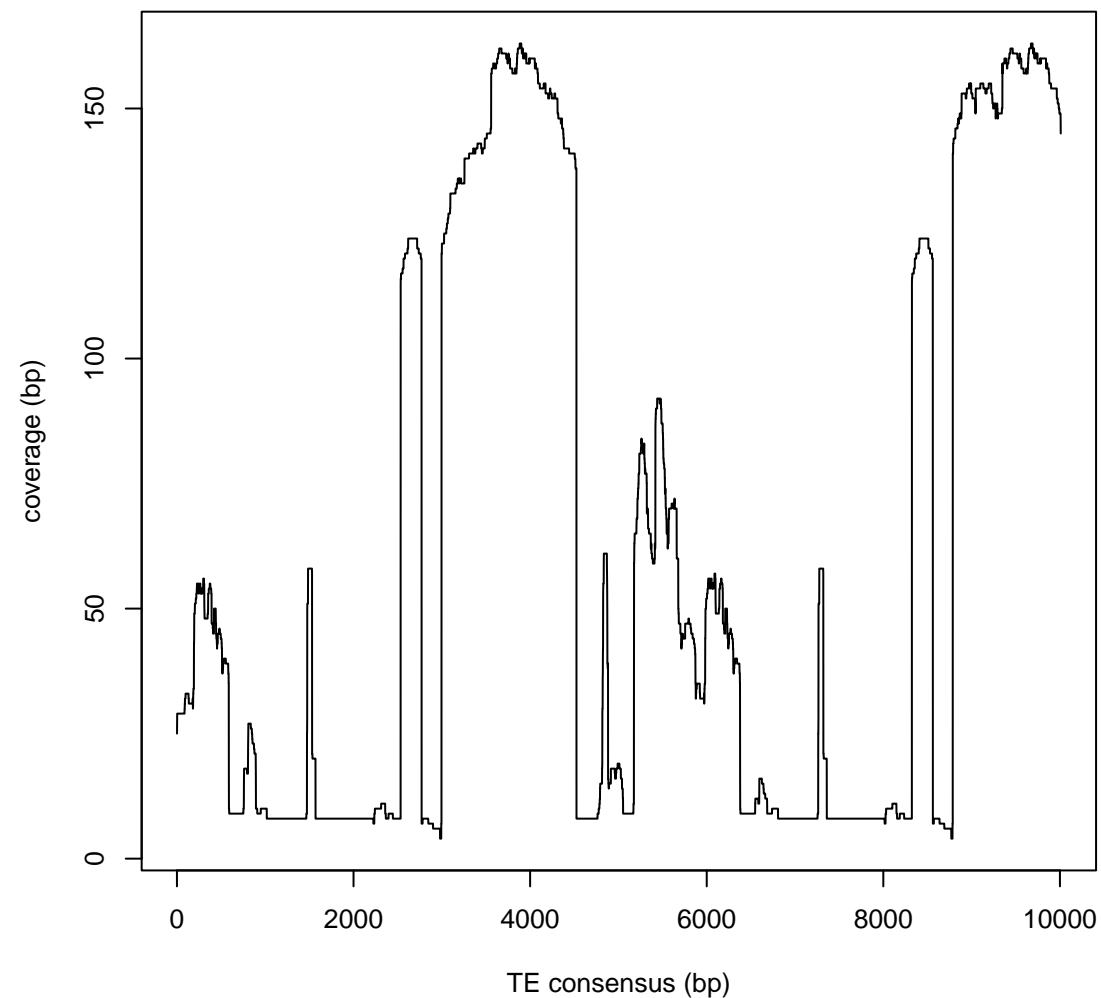


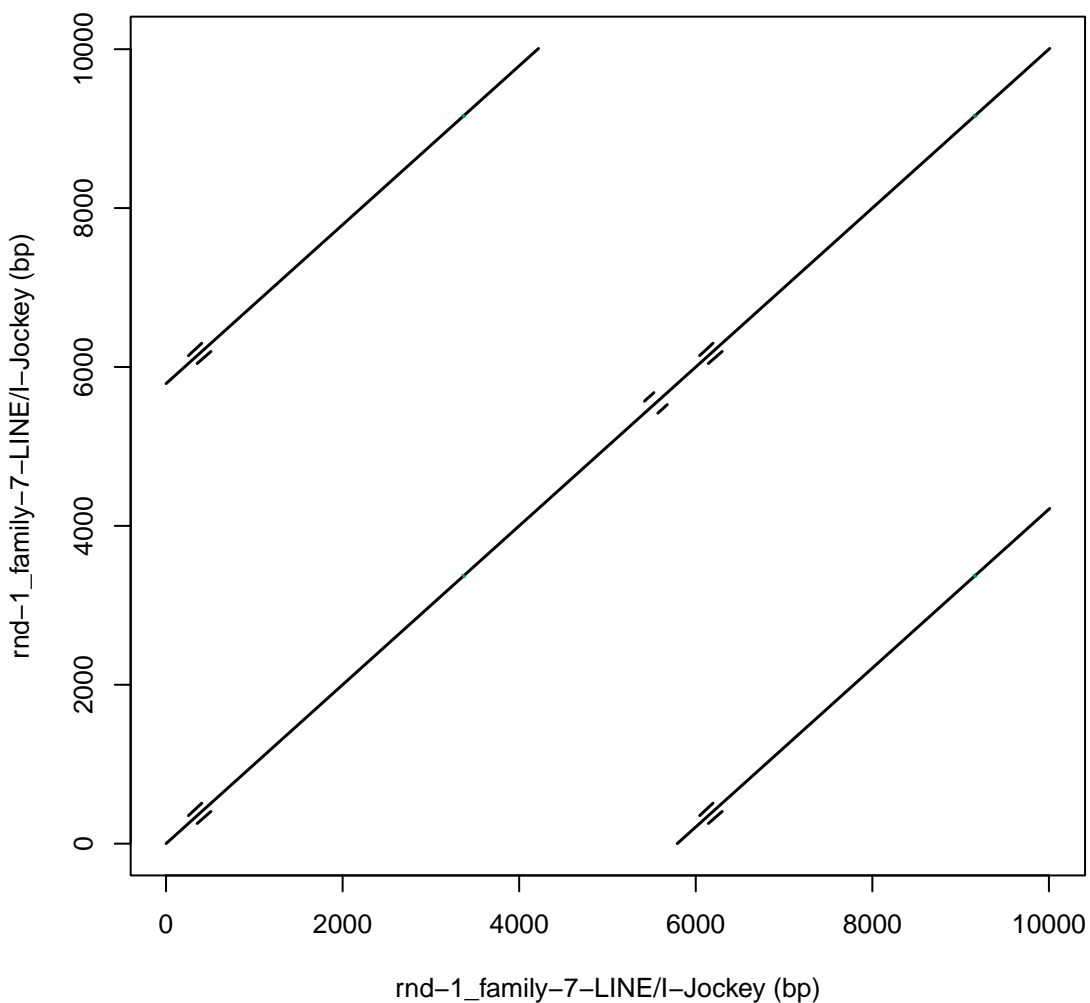
TE: rnd-1_family-7-LINE/I-Jockey
consensus size: 10009bp; fragments: 1109; full length: 2 (>=9008.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

