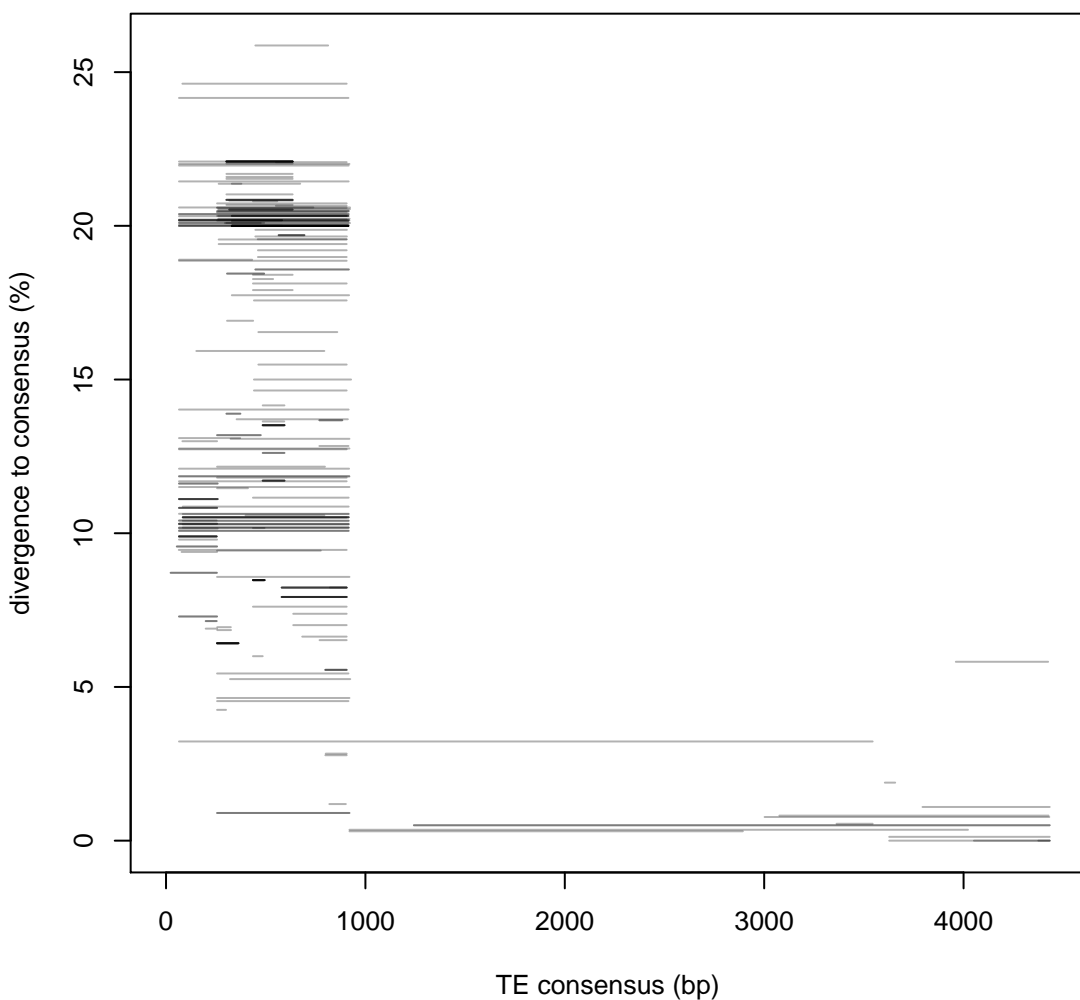
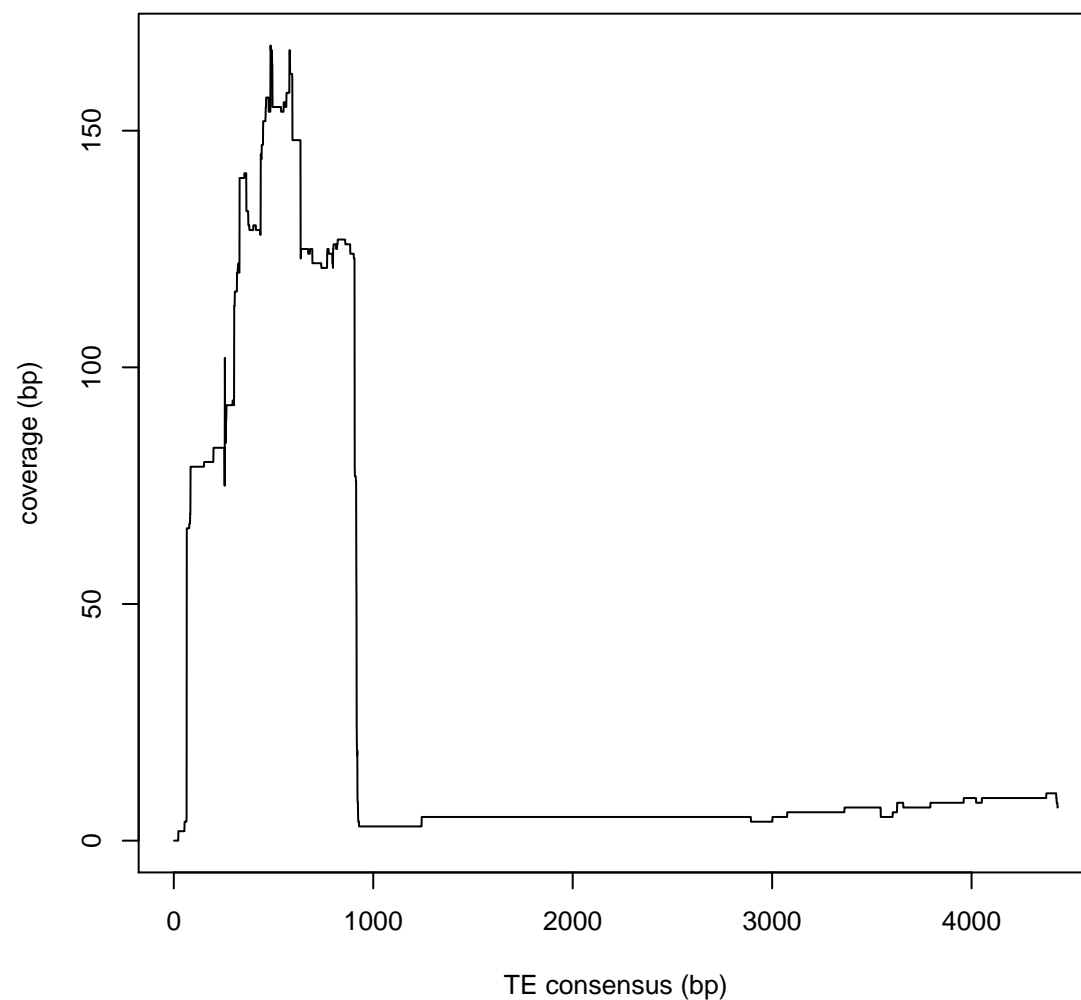


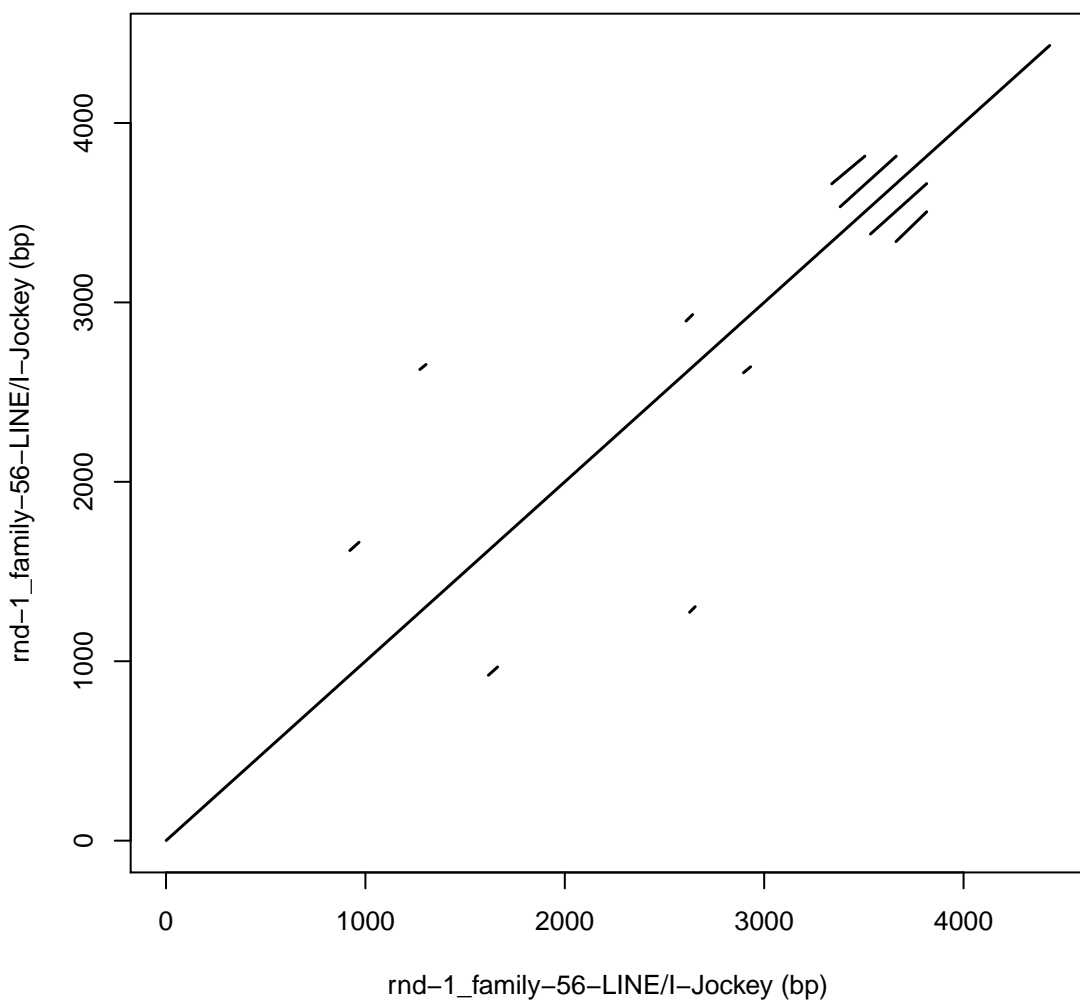
TE: rnd-1_family-56-LINE/I-Jockey
consensus size: 4432bp; fragments: 261; full length: 0 (>=3988.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

