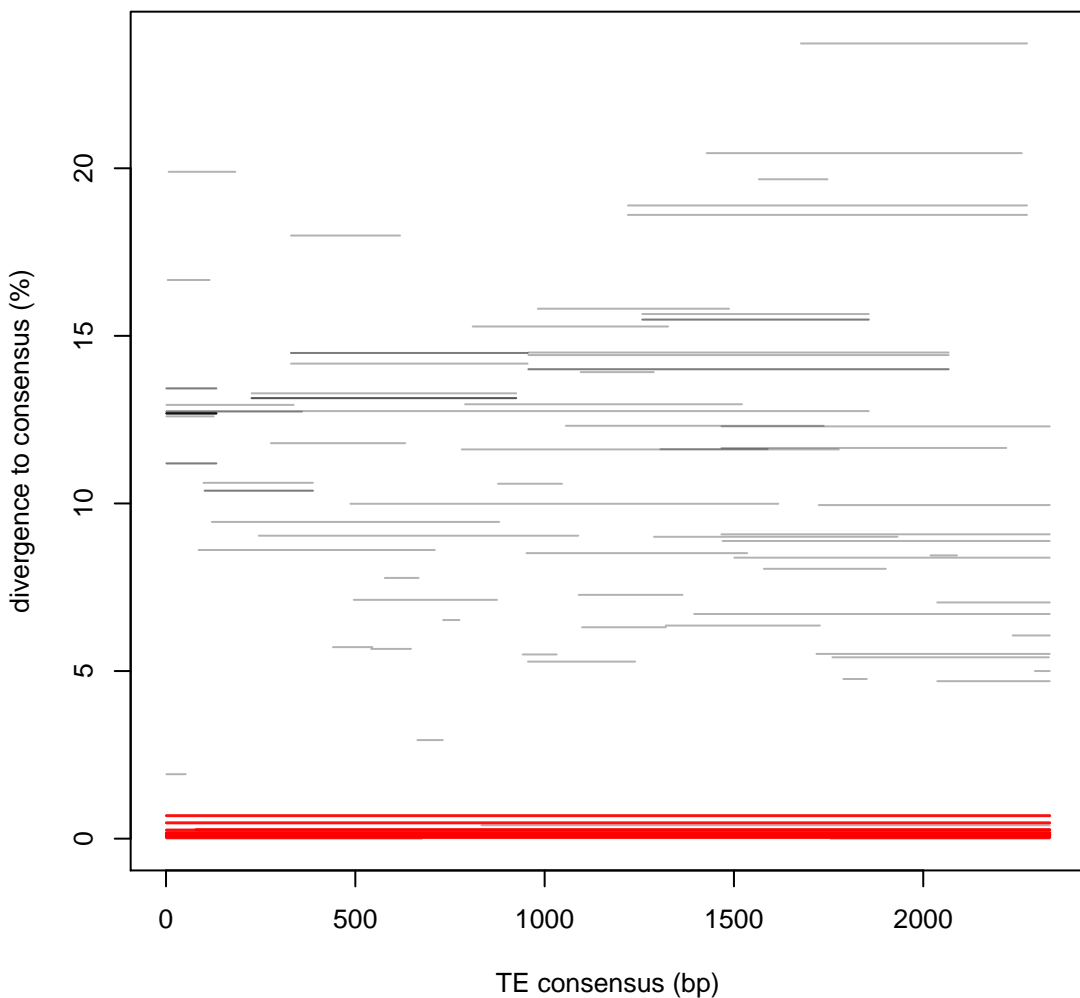
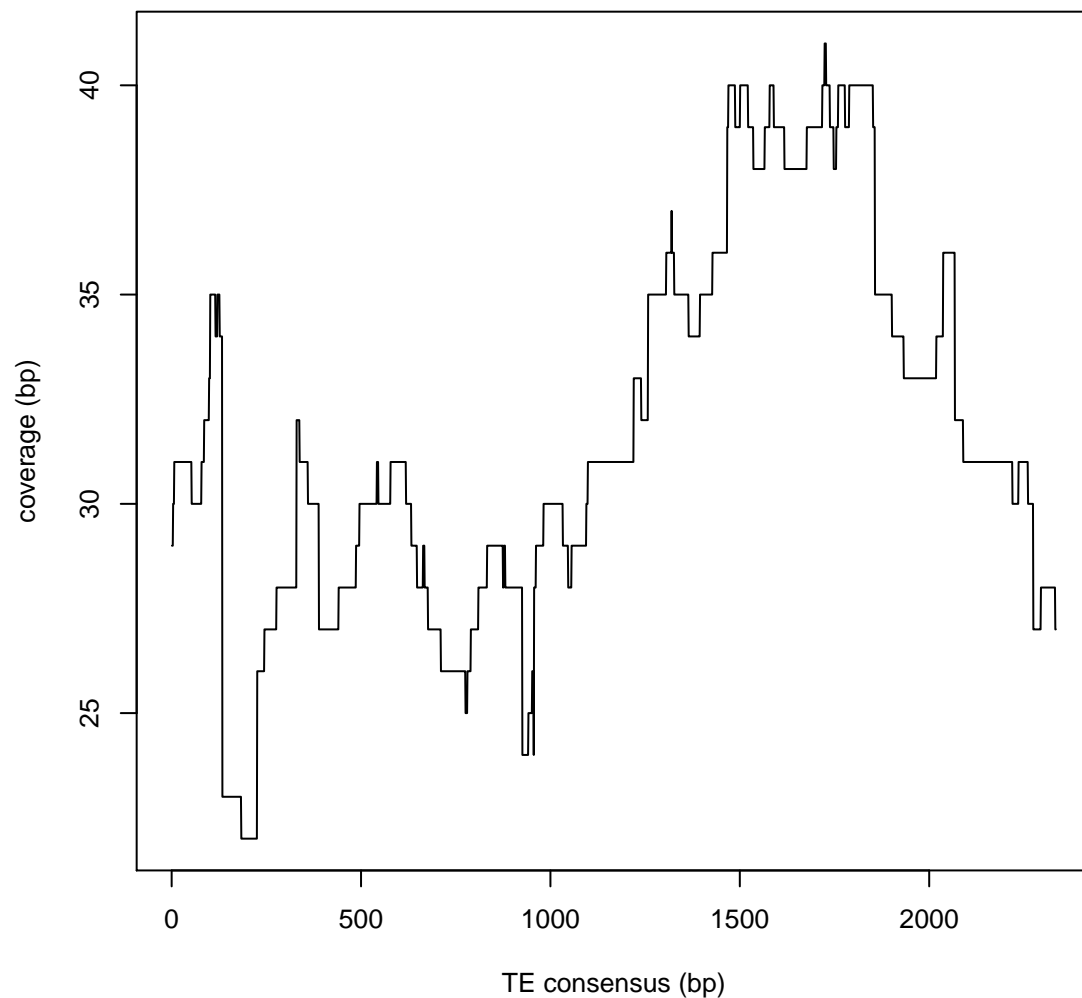


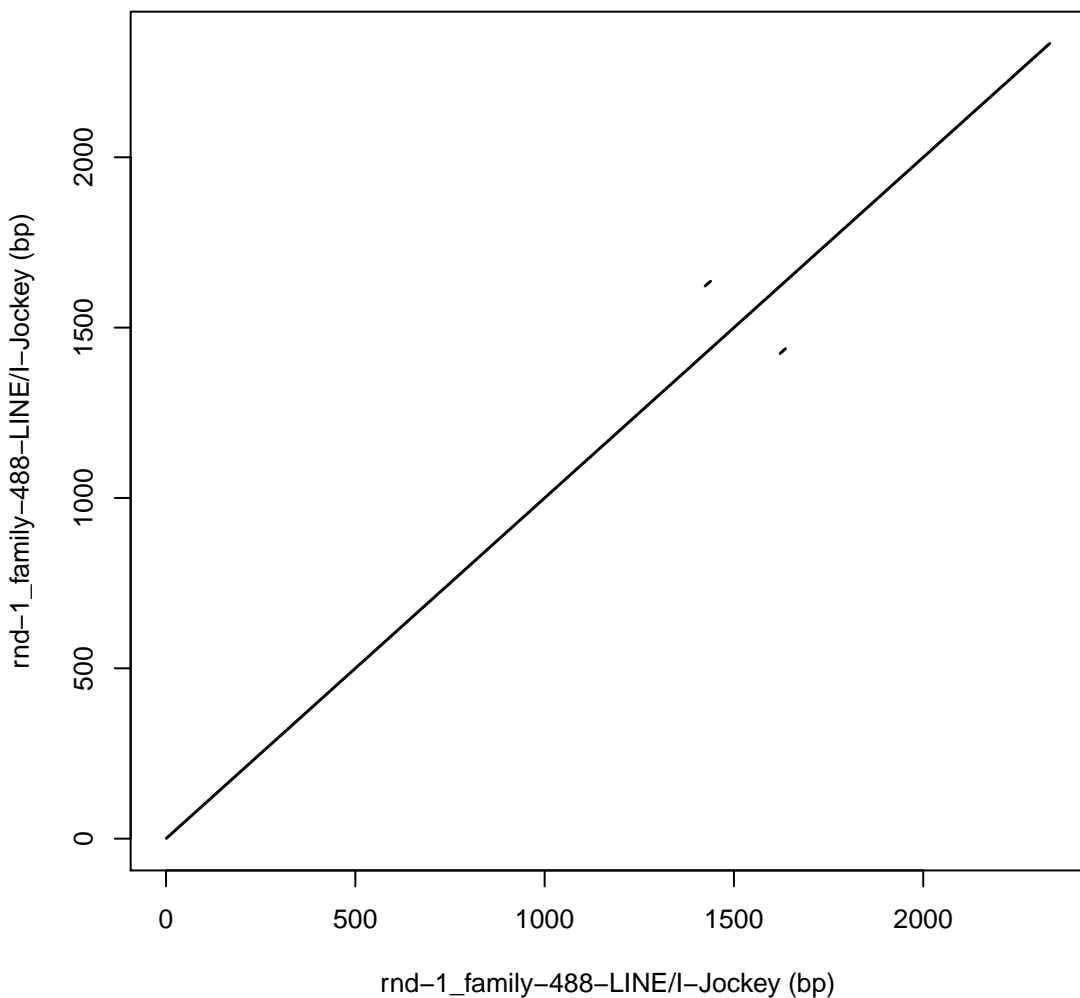
TE: rnd-1_family-488-LINE/I-Jockey
consensus size: 2334bp; fragments: 100; full length: 13 (>=2100.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

