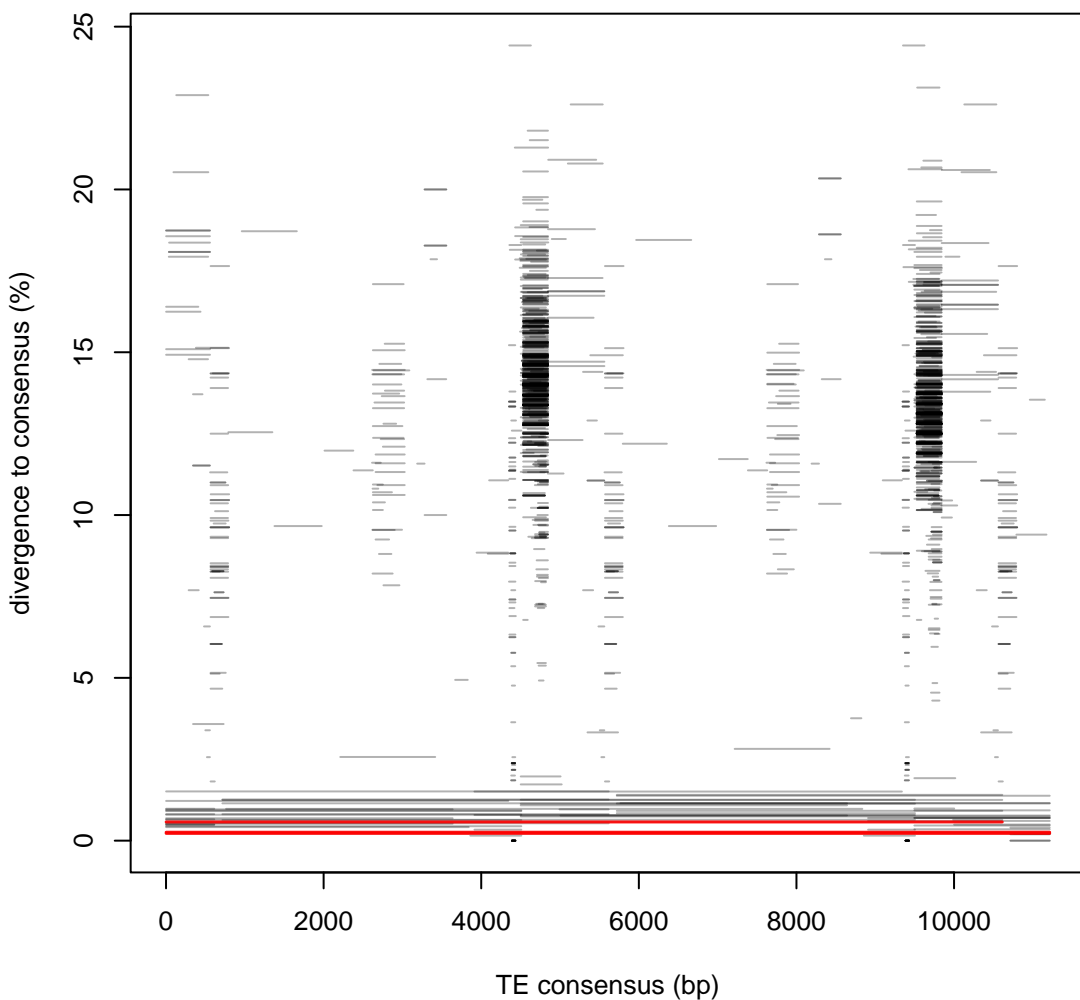
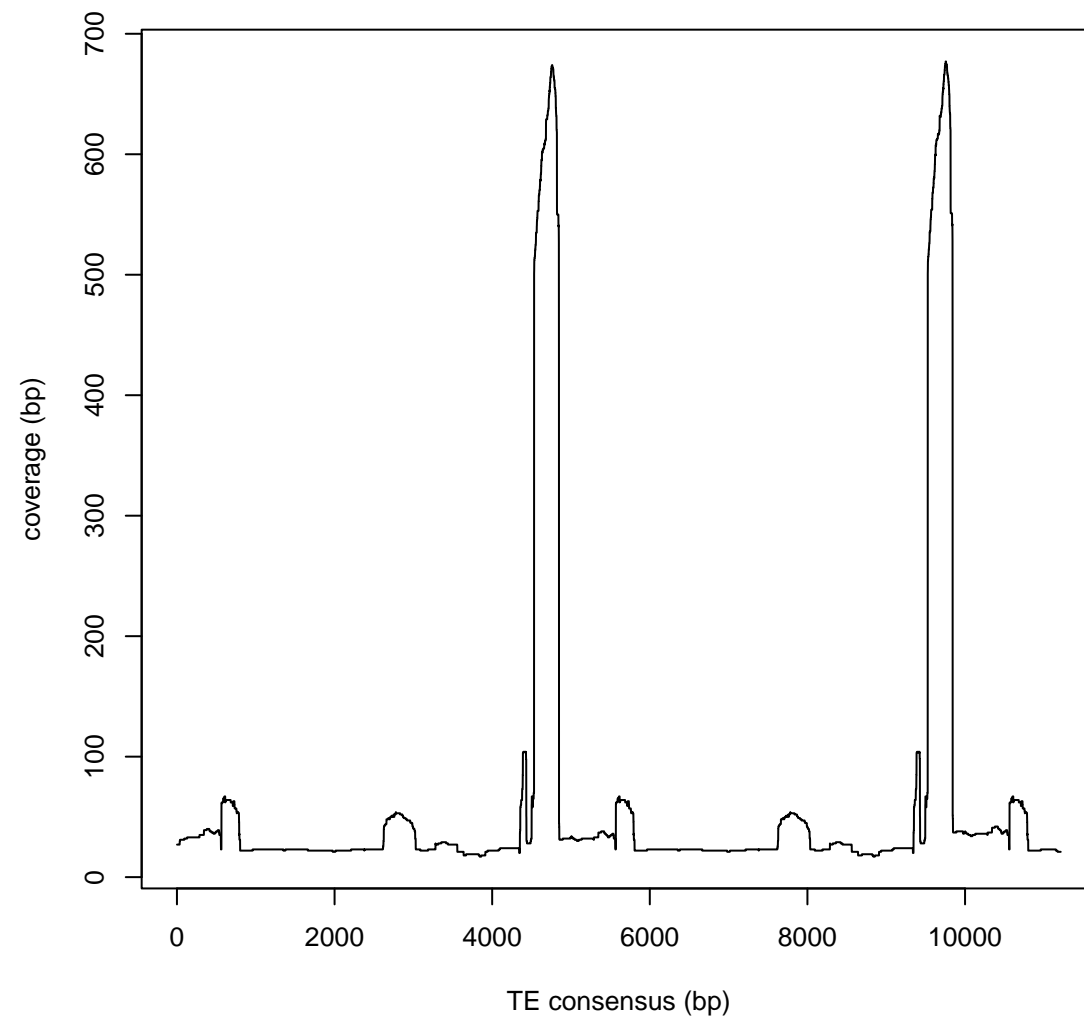


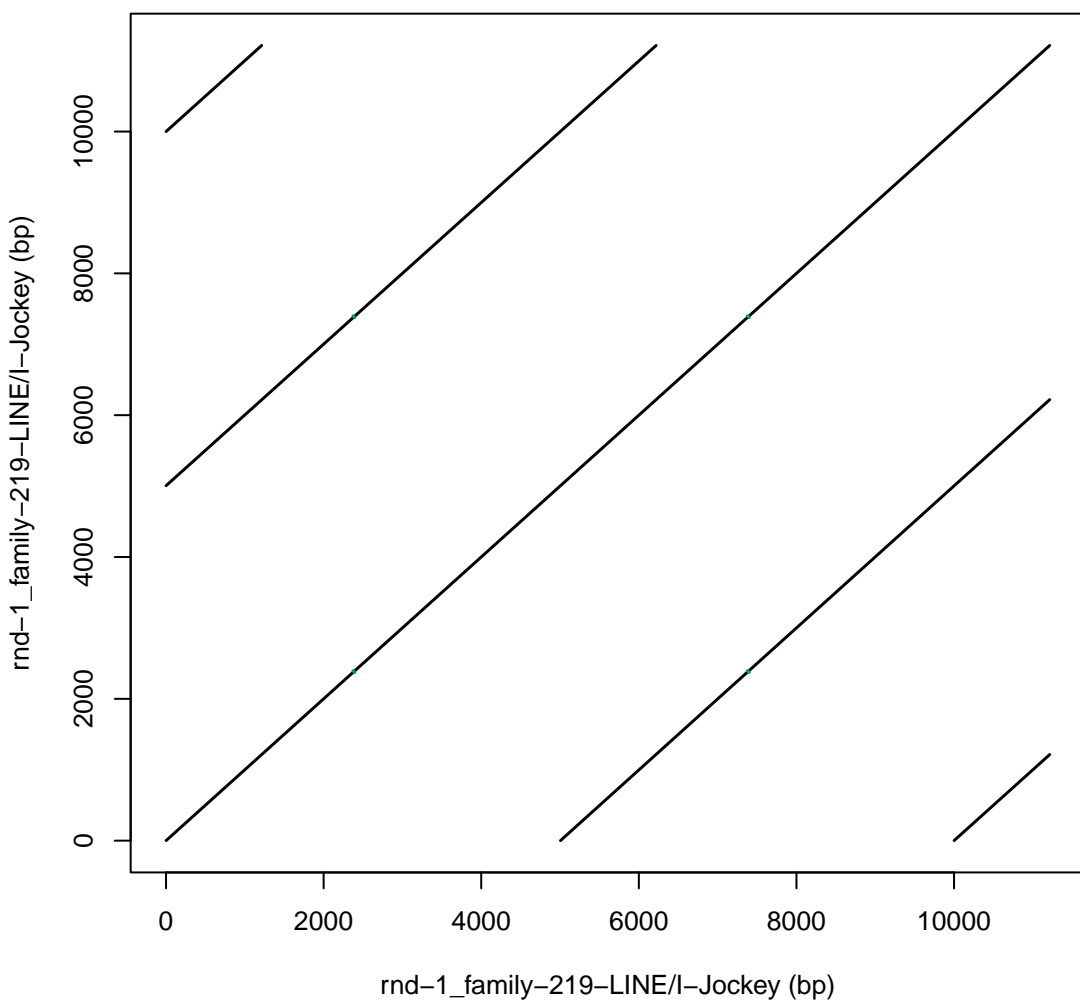
TE: rnd-1_family-219-LINE/I-Jockey
consensus size: 11214bp; fragments: 1887; full length: 3 (>=10092.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

