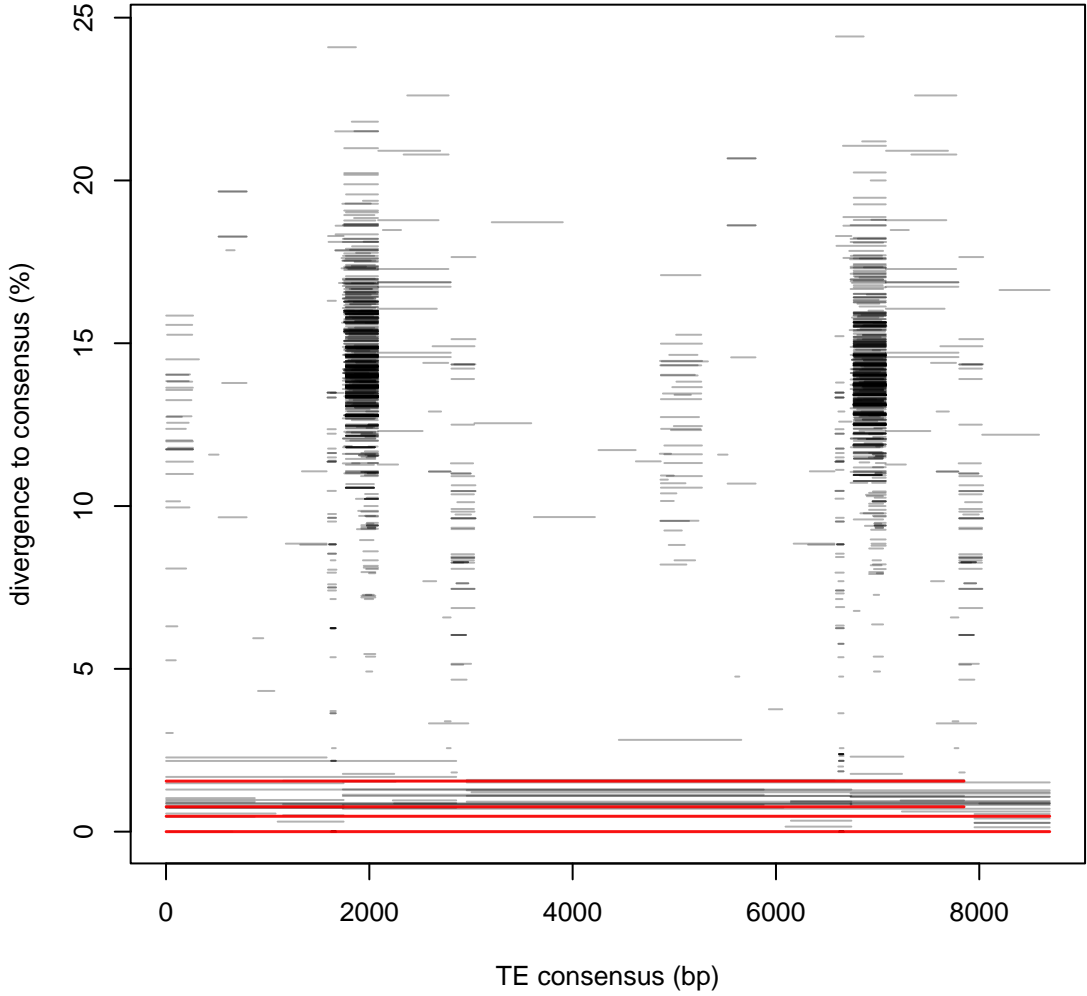
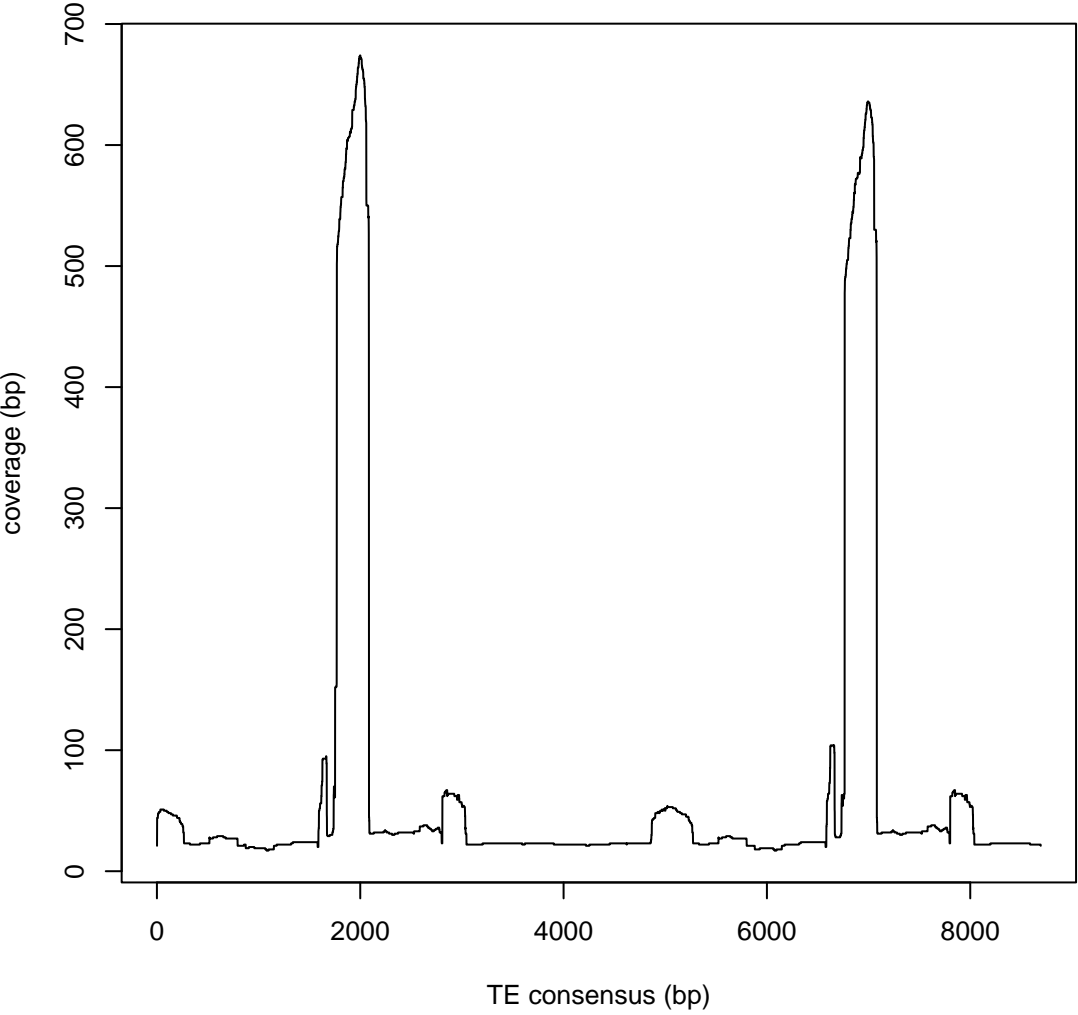


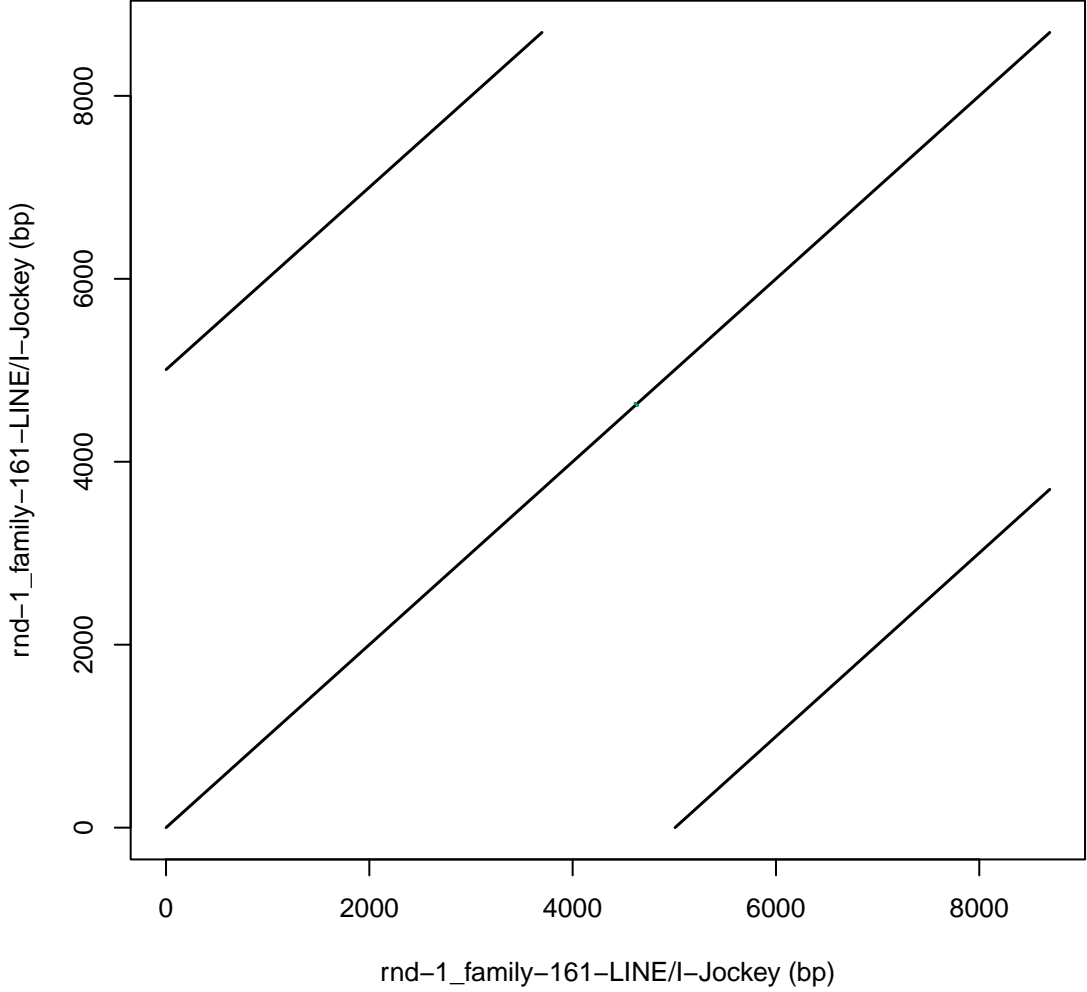
TE: rnd-1_family-161-LINE/I-Jockey
consensus size: 8693bp; fragments: 1741; full length: 4 (>=7823.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

